

GCTCAGGTTAGCCCGAAAGATGTTAGTATAGATTAGGCCACTCTAATTAAATGATCACTGAAAATACCTACTGGCATACTAA
CCAAGAGAATGCAAAAAAGCCGTCTCAAATTATGAAACAGCTATCAACAATTATAA

(SEQ. ID. NO. 195)

MEGLVRVHLLPVFGDYKLSKLTTPILOQQQVNWKADKANKGEKGAFANYSLLHNMNKRILKYGVAIQVIQYNPANDVIVPRKQQ
KEKAAVKYLDNKELKQFLDYLDALDQSNEYENLFDVVLYKTLLATGCRISEALALEWSIDLESGVISINKTLNRYQEINSPKS
SAGYRDIPIDKATLLLLKQYKNRQQIQSWKLGRSETVVFSVTEKYAYACNLRKRLNKHFDAAGVTNVSFHGFRHTTMMY
AQVSPKDVQYRLGHSNLIMITENTWHTNQENAKKAVSNYETAINNLZ --

Please delete the first paragraph located on page 101, line 1, and replace therewith the following replacement paragraph. A marked-up version showing changes made to the replacement paragraphs is contained herewith as Appendix B.

--What is claimed is:--

REMARKS

Claims 1-20 are currently pending in this application. Upon entry of the amendments contained herein, claims 1-22 will be pending. No new matter is introduced into the specification by way of these amendments. The Specification amendments are introduced to conform the application to the requirements of 37 C.F.R. § 1.77 and 1.821-1.825. Support for the claim amendments can be found, *inter alia*, throughout the application as originally filed, for example, in the originally filed claims; on page 9, last two full paragraphs; and in Table 3. Support for new claim 21 can be found, *inter alia*, throughout the specification, for example, on page 9, last two full paragraphs. Support for new claim 22 can be found, *inter alia*, throughout the Specification, for example, on page 9, last two full paragraphs and on page 13, third full paragraph.

Priority

The Examiner has made acknowledgement of Applicant's claim of foreign priority based on "German" foreign application number 9816337.1. Applicants respectfully state that the claim to foreign priority is in fact based upon Great Britain priority application number 9816337.1. Applicants respectfully direct the Examiner's attention to the executed declaration, claiming the benefit of GB 9816337.1, as well as the cover sheet of International Publication Number WO 00/06738 (reciting the priority application GB 9816337.1, and of which the instant application claims benefit), attached herewith as Appendix C. Applicants will provide the Examiner with a certified copy of Great Britain priority application GB 9816337.1 within a reasonable time of receipt of the document.

Restriction Requirement

In the Office Action dated September 22, 2002, the Examiner required restriction of the pending claims under 35 U.S.C. § 121 between one of the following inventions:

- “1.) Claims 7-8 and 13 drawn to DNA, host cell and a vaccine composition, classified in class 536, subclass 23.7. Further election of invention required.
- 2.) Claims 1-6 and 10-12 drawn to polypeptide, classified in class 424, subclass 190.1. Further election of invention required.
- 3.) Claims 15-16 drawn to an antibody classified in class 530, subclass 388.6. Further election of invention required.
- 4.) Claim 9 drawn to a method for inducing immune response using polypeptide classified in class 424, subclass 165.1. Further election of invention required.
- 5.) Claim 20 drawn to a method of use for the treatment or prophylaxis of *S. pneumoniae* infection using protein or polypeptide classified in class 424, subclass 184.1. Further election of invention required.
- 6.) Claim 19 drawn to a method of determining whether a protein is anti-microbial target in vivo or in vitro classified in class 424, subclass 93.1. Further election of invention required.
- 7.) Claim 18 drawn to a method of detecting *S. pneumoniae* infection using nucleic acid, SEQ ID NO:1 classified in class 435, subclass 6. Further election of invention required.
- 8.) Claim 14 drawn to a method of detecting *S. pneumoniae* infection using protein antigen, SEQ ID NO:2, classified in class 435, subclass 7.22. Further election of invention required.
- 9.) Claim 17 drawn to a method of detecting *S. pneumoniae* infection using antibody classified in class 435, subclass 7.22. Further election of invention required.”

See Paper No. 11, page 2. The Examiner further required election of one sequence falling within one of the above-identified inventions.

Applicants were contacted by the Examiner and invited to make an election to a restriction requirement by telephone. The Examiner originally designated (by telephone) as separate inventions the following 5 groups: (I) *S. pneumoniae* nucleic acid sequences (claims 7-8 and 18); (II) *S. pneumoniae* peptide sequences (claims 1-6 and 9-14); (III) antibodies to *S. pneumoniae* antigens (claims 15-17); (IV) methods of assay design (claim 19); and therapeutic methods (claim 20). Based upon this grouping of claims, Applicants provisionally elected by telephone on July 09, 2002, without traverse, the invention of Group V, claim 20, *drawn to therapeutic methods*. To the extent that the restriction requirement in the Office Action dated September 20, 2002, differs from that described in the telephone restriction to which Applicants responded on July 9, 2002, the restriction is traversed. In particular, Applicants submit that there is substantial overlap in scope between Group 4 and Group 5 or Group 6. Dependent claims to particular embodiments may properly fall within the scope of both generic claim 9 and generic claim 20. Likewise, dependent claims to other

embodiments may properly fall within the scope of both generic claim 9 and generic claim 19. Applicants respectfully request that the restriction requirement be recast.

Applicants also submit that the requirement for election of a single nucleotide (polypeptide) sequence is properly characterized as an election of species. Applicants provisionally elect the invention of claim 20, now classified in Group 5.

Consistent with the requirement of election of species, Applicants hereby further elect the protein sequence as set forth in SEQ ID NO:162.

Rejections

35 U.S.C. § 101

The Examiner rejected claim 20 under 35 U.S.C. § 101, as allegedly improperly defining a process. Applicants have amended claim 20 to conform with U.S. Rules of Practice, thereby rendering the Examiner's rejection moot. Accordingly, Applicants respectfully request reconsideration and withdrawal of the rejection of claim 20 under 35 U.S.C. § 101.

35 U.S.C. § 112, 2nd Paragraph

The Examiner rejected claim 20 under 35 U.S.C. § 112, 2nd Paragraph as allegedly indefinite for failing to particularly point out and distinctly claim the subject matter which Applicants regard as the invention. More specifically, the Examiner stated that "the claim does not set forth any steps involved in the method/process, [and therefore] it is unclear what method/process applicant is intending to encompass." *See Paper No. 11, Page 7, lines 4-6.* Similarly, the Examiner stated that the claim is further indefinite because "the claims do not recite positive active steps so that the claims will set out and circumscribe particular area with reasonable degree of precision and particularly [sic] and make clear what subject matter the claims encompass, as well as make clear the subject matter from which others would be precluded." *See Paper No. 11, Page 7, lines 9-13.*

The Examiner further determined that "additional structural parameters, such as amino acid sequence of the polypeptide (for example; SEQ.ID.NO), would enable more definitive identification of the claimed polypeptide without ambiguity." *See Paper No. 11, Page 7, lines 17-19.*

The Examiner also rejected claim 20 as allegedly vague for the recitation of "capable of" within the claim, as this recitation allegedly makes it difficult to determine the metes and bounds of the claim.

Applicants have amended claim 20 to conform to U.S. practice. As a result of the claim amendment, the Examiner's rejection of claim 20 has been rendered moot. Applicants respectfully request withdrawal of the rejection of claim 20 under 35 U.S.C. § 112, 2nd Paragraph, and reconsideration of the application in view of the amendments.

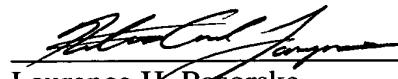
Conclusion

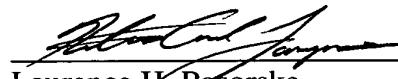
Applicants believe that amendment of the subject application has placed this application in a condition for allowance. Early notification of a favorable consideration is respectfully requested.

Respectfully submitted,

Dated: March 20, 2003

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IN THE UNITED STATES PATENT AND TRADEMARK OFFICE

In re Patent Application of :)
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)
 GILBERT, C.F.G., et al.) Examiner: Baskar, P.
)
Serial No.: 09/769,787) Group Art Unit: 1645
)
Filed: January 26, 2001)

For: NUCLEIC ACIDS AND PROTEINS FROM STREPTOCOCCUS PNEUMONIAE

Under Secretary of Commerce for Intellectual Property and
Director of the United States Patent and Trademark Office

Washington, D.C. 20231

Appendix A - Version with Markings to show changes made to the Claims

In claim 20:

--20. (Amended) A method for treatment or prophylaxis of *S. pneumoniae* infection comprising administering to a patient in need thereof The use of an agent capable of antagonizing, inhibiting or otherwise interfering with the function or expression of a protein or polypeptide as defined in Tables 2-4 in the manufacture of a medicament for use in the treatment or prophylaxis of *S. pneumoniae* infection, wherein said agent induces an immune response in said patient to a polypeptide having a sequence comprising SEQ. ID NO:162.--

IN THE UNITED STATES PATENT AND TRADEMARK OFFICE

In re Patent Application of :

GILBERT, C.F.G., *et al.*

Serial No.: 09/769,787

Filed: January 26, 2001

) Examiner: Baskar, P.

) Group Art Unit: 1645

For: NUCLEIC ACIDS AND PROTEINS FROM STREPTOCOCCUS PNEUMONIAE

Under Secretary of Commerce for Intellectual Property and
Director of the United States Patent and Trademark Office

Washington, D.C. 20231

Appendix B - Version with Markings to show changes made

In the paragraphs starting at page 29, 1st full paragraph (immediately below the heading "Table 1") and extending to and including the last full paragraph located on page 100:

4101.1 (SEQ. ID. NO. 208)

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4101.3 (SEQ. ID. NO. 209)

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GGAA

4101.5 (SEQ_ID_NO. 210)

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4102.1 (SEQ. ID. NO. 211)

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4106.1 (SEQ. ID. NO. 212)

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4106.4 (SEQ. ID. NO. 213)

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4106.6 (SEQ. ID. NO. 214)

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CGTCAAGGCTTAATGGAAGAAACAGGATTCTCAAAGCAACCTAACCAAATATGTCACCCCTGCTCAATGACAAGGTTG
GATAGTGGCTTAGAGCTGGCTATTCACTCAGAAGATGAAAATCTGGCTGTCTATCGGTGAGCTACCAAGGGAGAGATA
TTGGAGCTTGTGTTGGAGAGTGTGTTAATACAGATTGGTTATCTCTTCTACCCACAAAGTTTACGGCCATCAGC
TGCTCAAGAATTGGTATTAGCAGGCTACGTTGGTCGTACTTGGTGGTTAAATCAGATTGTCAGAATTGATT
TCCATCCAAAATGGCGTTGGCGAGGTCCAGAGCATCAGATTCACTTCTATTCTGCTTCTGGCAAGGCTGGTCA
TCAGGAATGGGAAGGTCACATGCAAGAAACAGAGAGAAAACAGGAGATTGCAAAATTAGAGGAAATCTGCGGTGCAAGTT
GTCTGCGGGCAGAAATTGACTTGTCTGGGCTACATCAGTCAACACGCTTCTGGGCTCAATTGCTGAGTTCAAG
TCATAGAAGAGAAAATGCGAGGGTATTGACAATTCTTCTGAGGAAAGGTTCCGTCCTTGTGG
CACACATATTCCACTAGGAGTTGAGGATGGTGAAGATGATGATATTCTCTTCTCTATCTCATCGATTCTCTTCTCAT

ACTATGGAGTATATTCTGGTTTGAGGGCAGTTGGCAGATTACTGACGAATTGATTCAAGAAATGAAGAAGGGAGGAAC
TATTGGGGATTATAACAGAGGACCATGTCACCTATGAACACTCAGTCAGCTTGTGCTCAAGTCTATCTCTATAAGGGCTATATT
TTACAGGATCGCTACAAGTACCACTAGAGAATCGTCATCCATATTACTGATGAAACATGATTAAAGAGACAGCAGAGG
AGATTTTCTATGCTCACCTGCTTCAACAGGGGACAGATTAGATAAGAAGATTCTCTGGGATGGCTCCAGTTAACGAA
TATATGGCTGAAAACGGTGGGCCAGCATATGCGGATTGCTGGATITGACATCTGGTCTTCAAGGATGGCAGC
CATTITGAAACGGTATTGGAAACATACGTTTATTACCATTAAGCCTATGACCTAGTCGGCATTATGATTGCTGGTTAC
CAAATAACCCGATTCTATAAGAAGGAACAGACACCAGTCTATTATITAAAAATGACTGGATATGGAGGATITGGTAGCGATT
CGCCAGTTATTACTAA

4106.7 (SEQ_ID_NO. 215)

ATGGAATTTCAAAAGAAAACACGTGATTGCAATTAAAAAAATGCAGGAACGTACCCCTGGACCTCTGATTATCGTGGAG
GAATCACAGGAGCTGGTGTAGCTTGCAGGGCAGCTAGCGGTCTTGAGACTGGTTGATTGAAATGCAAGACTTGCAGA
AGGAACATCTAGTCGTCACAAAATTGGTCACGGAGACTTCGTTACCTAAACAAATTGACGTTAGAAGTGGTCTCAGAT
ACGGTTCTGAACGTGCACTGGTCAACAAATCGCTCCACACATTCCAAAATCAGATCCAATGCTCTTACAGTTACGATGA
AGATGGAGAACCTTAAAGCTTCTCGTCTTAAAGTAGCCATGGACTTGTACGACCTTGGCAGGTTAGCAACACCCAG
CTGCGAACAGGTTTGGAGCAAGGATCAAGTCTTGGAACGCCAGCCAAACTTGAAGAAGAAGGCTGGTAGGAGGTTGAG
TGATCTTGTACTCCGTAACAGATGGCGTCTCGTGTGATTGAAAACATCAAACGTCGCAACCAAGACGGTCCCCATTGCC
AACACGTGAAGGCAGAAGGCTTCTTGTGACGAAAGTGGCAAGATTACAGGTGTTGAGCTCGTGTACTTGTGACAGACC
AAGTGGTGAATCAAGGCCGCTGGTTATTAAACACAGGTCTTGGAGTGTAAAGTACGTAATTGTCTAATAAGGG
AACGCAATTCTACAAATGCGCCCAACTAAGGGAGTTCACTTGGTAGTAGATTCAAGCAAATCAAGGTTACAGCAGTT
TACTTCGACACAGGTTGGTGGAGCTGCTATGGTCTTCTCCACGTGAAACAAAGACTTACTTGTACAACGTGATAC
AGACTACACAGGTATTGGAGCTCAAAGAAACTCAAGAAGATGTAGATTACTACTTGTGCAACAAACCGCTTC
CCGAAATCAAACATCACCATTGATGATATGCAAGCAGCTGGCAGGTCTTGC
TC CATTGATTGCAAGGAAACAGTGCCTCTGACTATAATGGTGGAAATAACGGTACCATCAGTGATGAAAGCTTGTACAACITG
ATTCGACTGTTGAATCTTATCTCTCAAAGAAAAACACGTGAAGATGTTGAGTCTGCTGTGCAAGCTTGTGAAAGTAGCA
CATCTGAGAAACATTGATCCATCTGCACTTCTCGTGGGTCTAGCTGGACCGTGTGACAAATGGTCTTGTGACTTGT
GGTGGTAAACACAGACTACCGTAAGAGTGGCTGAAGGAGACTGGAGCGCTGGACATCTCAAAGCAGAAATTGAC
CGTAGCTTAAATTGATCAATTCTAAACTTACCCCTGTTAGGTGGAGTACGAAAGGAAGCTCACTATCTGGCAAATCTTACGGTCAATG
AACGAAAGTCTTGCACTTGCACAGCTTGGAACAGGCCAGGACTCAGCTGGCAGATACTTGTCCCTCACTATGCAA
TGCACATGAGTTGACTCTAGCCCAGTTGACTTCTTCTCGTGTACCAATCATGCTCTTATGCGTGTAGCTGGATA
GTATCGTGTAGCCAATTGGATGAAATGGGACGATTCTATGACTGGACAGAAGAAGAAAAGCAACTTACCGTGTAGTGT
CGAAGCAGCTCGCTAACACGATTAGCAGAAATTAAAAATTAA

4106.8 (SEQ_ID_NO. 216)

ATGATGAATGAATTATTGGAGAATTCTAGGGACTTAACTCTGATTCTTAGGAAATGGTGTGAGGTGTGTTCT
TCCTAAACCAAGAGCAATAGCTCAGGTTGGATTGATTACTATGGTTGGGGATTGCACTTGCAGTTGGCTGAGCTTGTAT
CTGGCAAGCTCAGTCCAGCTTAAACCCAGCTGTGACCATCGGTCTGGCTTAAAGGTGTTGGCTGACATTCAAACCTCACTATG
TTGCTTATATCTTACGCCCCAGTTCGAGGGCCATGCTGGTCTAGATTGGTGTGCAATTCAAACCTCACTATG
GGTGGTAAACACAGACTACCGTAAGAGTGGCTGAAGGAGCTGGAGCGTGTGACATCTCAAAGCAGAAATTGAC
CGTAGCTTAAATTGATCAATTCTAAACTTACCCCTGTTAGGTGGAGTACGAAAGGAAGCTCACTATCTGGCAAATCTTACGGTCAATG
AACGAAAGTCTTGCACTTGCACAGCTTGGAACAGGCCAGGACTCAGCTGGCAGATACTTGTCCCTCACTATGCAA
TGCACATGAGTTGACTCTAGCCCAGTTGACTTCTTCTCGTGTACCAATCATGCTCTTATGCGTGTAGCTGGATA
GTATCGTGTAGCCAATTGGATGAAATGGGACGATTCTATGACTGGACAGAAGAAGAAAAGCAACTTACCGTGTAGTGT
TCGGAGCAGCTTGGCAGTGTGTTGATTCTACCTTCTAG

4106.10 (SEQ_ID_NO. 217)

ATGAAAAGGACCTGGAGGAACCTTCGACAAATCTAATACACCTTTATGATTGCAATATTGAGATTCCAATCGTAC
CGTTTACGGCCTATGGCTGGCGTGCACCAACTCAGCCTTCTGACTATCGCAAGAGCTGGAGCTGGACTCGTTGTAATG
AAATGGTCTGACAAGGAAATCCAATACAACAAACGAAAACCCCTGCACATGCTCATATCGATGAGGGCGAAAACCCCTG
TCTCTATCCAACCTTGTGAGCATGAAAGACAGCCTAGCACCGCAGCAGAAATTCTACAAAGAAAACCCAAGACCGATAT
CGTCGATATCAACATGGCTGCCCTGTCAACAAAATCGTAAGAACAGCTGGCTATGGCTCAAGGATCCAGACAAG
ATTACTCCATCATCAACAAGGTCAGTGTCTTGATATCCCACCTACTGTCAAAATGCGTACGGCTGGCGACCCATC
TCTTGAGTAAAGGACTCTCGTGTGAGACTGCAAGGCTGGCTGCCATGCGTACGGCTACCCGTGAAACAAATGT
ATACTGGCCACCGCAGACCTTACAAGGGTGGCCAAGCTCAACCAAGTCCATCGCCAACGGTGTGATAC
CGTACTTCAAGAACGCAAGAACGCAACGCTGAGAAGATGGTGTGACGAGCTGATGTTGGCGAGCTGCCATGGGAAAT
CCCTACCTCTCAACAAATCAACCAATTGAAACGATTGATTAACCTCAAAGGAGAAAACGTCGAGTCGTAATTCCGCGTCTCGTCTCAC
TATCTCCGTGAAACATCTGGCGTGCCTAACCTCGTGGAGCCATTGCAAGCCAGCACCTGGCAGAGATTGAAACCCCTCT
TGCAATTGGAGAAGGCTAA

4107.1 (SEQ_ID_NO. 218)

ATGACAAAGAAGAAAATTGAGCGTATTCTGTAATACACCGAGAAAAGATTATGGCTCAAGTGGTATTCTATGCGAGATA
AAAACAACTCAAGTATAGTGTCTTGAGCGTAAATGTTGATGCTGCTAAACAAAGATATGCTAGCTTATCAAAAATA
CGCAACTATCAAGCAGATAACAGATATTAGGTACAAACAAAGTGGCTGACATTAGGGCTGAAAGAGGTTATGTG
TACAATCACATGAATGTTATCGGAGCTGTGACGGATATTATTCAGTCAATCACCAGCTTATGATAAGTTAATAAGTG
GTTAATATCTATTCTGATTGTTAGCGTTGACCTTGCCTAAAGGGGTATATCATGAGATGGTAGGTATCTAG

4107.2 (SEQ_ID_NO. 219)

ATGAAAAGTCCAAACGGGCTGAGATGAAATTACTTTATACTGATATTGGACTTCTTGCAGAGAAATTCTAACAAAGAGG
CGAAGAGCTAGTTGCACTGGCAAGGGCTTCTACATTGCCCAACTCTTCTTCTTCTGAAAAGGAACGGCCCGTCTG
GAATACTTGTCCAGCAGGCTTCTTCTGATTACCGTCACCGCTTGTCAAAATGGCTCGTATCTGGTCTTGAATGATT
CCAGCTAAACTACTTGTGATGATATCGGTCTTGGGTTGGCTTACAAATGCCGACTCGATCCAAGGACTTGC

TGTTTATGGCGCTATTAAAGCAGGATCCTCAATTGATCCAGCAGTTAATTGAGCTTACCATGAGATGACCAAATCTCAGATGA
GTTTTTGAGCTTGGAGAATTAAACAGATGAGGATAAGAGGGCGGATTACTCTGATTTTGAGAAAGTAACAGCCTATCTT
AATCAAGGTCAGTTAGGCCAGGAAGTCAGTTGTCCTTACCTGAGGCTATTGAGAATGACAAGGTAAGTAGTGTATT
ATCAAATCGCTTGGTCAATTGACGGCTTACTCTGTTTCTGCTGAGGAAGAGCGGGTTGAGGACTTACCTCACGGCAAGG
GTTGAGATTGTTATCGGGCTTACTGAGTAAGAAAGCCTATACCAGCTTAACTGAGGACTGTTCTCAAACATGAGAAGATGG
AAAATTCTCATCTGGCTTCAAATACCAACGCCGCTCAGGACTGTTCTCAAACATGAGAAGATGGATAGTTG
ACAAGGCCTCTGTTGGAGTCTTATGACTTTAGAAGACTCGCTTGGATGTCGATGAGAAGAGCCGTAAAATT
CAAATCTGGCTTGGAGGAGCTTGGAGCTAGTAGGCCGAGTATTCTGAGAATTACATGAGAAGACTCAG
ACTGAGCTACAAGCATTCTGATTTCTGTTGGGGAGTAGCTGAGCTTCTTACCGATTCTCTCAAACACATTGAGGACT
AGATTCCTTTTATCTGGAGAAGCCGAGGCTCATCTGACTCATCCCTGACTGTTGAGCTTACCTGAGTCTTAAAC
GTTACCGTTTCGTCAGGGAGATTGAGTAATCTCTTAACTGAGCTTACTGACCTCAGTCTGAGTATTGAGCT
TTGAGCAATATATCCGCTATCTGGTATCAATGGCTTGCAGCCTTCAAGAACCTTACCAAATCCCACCATGGAAAATT
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AAAACCTCTACAAAAATGGAGTGTCTTCTAAAGAAGGAGCTGAGCAAGCAGTACAAGATTGACAACCACCTTGGAA
GCTGAGAAGCCGAAAGACAAGCCGAGTTGGAGGCTTCTGCTCATTGGAATGAGTTGGCCTTAAACCTGAGTCT
ACAGGTTAGCTGGAGAGACTCTGCTCATTGCTCATTGGAATGAGTTGGCCTTAAACCTGAGTCT
TTGAGCAATATATCCGCTATCTGGTATCAATGGCTTGCAGCCTTCAAGAACCTTACCAAATCCCACCATGGAAAATT
AATCTTGAGCGTTGAATGTCCTCCGAGGAGGAGCTTACCCACCATGAGCAAGACCTAGCCAAGGACACCTGAA
GCTCT
CTATCCTGCCAACAGGGAGTTTACCTGTCAGTCGGGTTGACAGAGTTTATCGCAATGAATACAGTTATTCCTACGCT
ACGTTTAGGCTTGCAGGGAGATTACGTTGACCTGAGGAACTTCTGCTGAGTCACGGGAAATTCTGCACTGCTATCTTGAACGC
GCCCTACAGTGCCTAATGAAGATTCTTGGACCAACGCTAGAAGAACCTATTCAAGAAACAGTCAGAACACGCAATTG
AAAGCTATTATCAAGAAAGTTGGAGGCCAGTTACCAAGGAAGTCTTGGGAAAGACCAAGCCTTATTCAATTAGACAAT
GGACGAGTGTCTTGTACGAGGGCAAGGTGGACCGGATTGACCGTTGAGCTAATGGAGCGATAGGAGTAGACTAC
AAATCCAGTCTGACTCAGTTCCAGTTCTCATTTAATGGCTCAATTCTGAGTACCTATCTGCTGCCCTAAA
AGAGAAGGGGAGCAGAACTTTGGCCCATGACTTGGAAATGGCTCAACTGAGTACCTATCTGCTGCCCTAAA
TGCCAGGAGCTGGTAGAAGCCAGAAACTCTGAGGAACTTCTGAGTACCTGAGTACCTGAGTACCTGCTGCCCTAAA
AATTTATAACAAAACAGGCTAATCAACTGACAGATGAGGAATTCTGAGTACCTGAGTACCTGAGTACCTGCTGCCCTAAA
GAAAGCTGCTGAGAAGATTAGCAGGCCGTTGCCATCAATCTTAACTGAAATGGAGCAGTACCTGAGTACCTGCTGCCCTAAA
CAGCAACATCAGGCTATTACAGGCTTGAAGCCAATTACCATCTGGCCAAGGCCGTTCTAGAAAAGTGGACCTAGCTG
ATGGCAAGGCTGGTGGAGAAAAACTCAAGCAAGCTTGGCTTGAAGAAAATT
GAGAGGAGTTGAATCGATGA

4107.3 (SEQ. ID. NO. 220)

ATGAAGCTTATCCCTTTAAGTGGAGGAGGAGTCAAAAAGTCAAGAAGCAGAAGCAAATCGAGCAAGGAACAGAAG
AAAACGCGAGCAAATCGAAGCTATCTACACTCTGCCAGAATATCTGGTCTCAGCATCGGCTGGITCTGAAAGACCT
TTGTCATGGCAGAGCGCATTCTGGCAGGAACTTGGCGCTGGTGCAGAATTCTCAACTCTTATCTCAACCTTACCGTCAAG
GCTGCACTGAACTTAAAGAACGTTAGAGAAAAAAATCAGCAAGAAAATCCAAGAAACAGATGATGTCGACCTAACAA
CACTGGGTCGCCAGTGGCAGACCTACCCAACGCTGCCATTGAAACCATGGATTCTTACACAAAAATCTTGGCAAAC
ATGGTTATCTGTTGATATTGACCTAATTCCGTTATTACAAACCAAGCGAGCAACTTATTCTGAAACAGAAGTCTT
CATGAGGCTTGAAGCGCATTACCAAGGAAACAGAGACCTTACTGCTGAAAGAACCTTGTGGCTGGCA
AGGACGACACGGGAGCTGGTAGAAGCCAGAAACTCTGAGGAACTTCTCCAAACGGCTTCTGCAACTCTGAG
TGAAATTTCTCTCAAAGGATTGAGAAAGCTGTTTACCATCTGAGTACCTGAGGAAACCTGAGTACCTGAG
TGGAATTGGAAAGCTTTCCGTTACCATCTGAGTACCTGAGGAAACCTGAGTACCTGAGTACCTGAG
GTTAATTCTGGATGAAATTGGCTCCCTAAATCAGGAGTCCGATAGTCAGGCTTATCAGGAGTCTGCTGCGCTGTTGCGCA
CTCTAAGGAGAAAACGGCTGAGGCTGACTAATGCCAGCGTAAGGCTGATTGAGGCCCTGGCTGATGCCCTAACACGA
AGAGGAGAAAAGACCCAGTTGCTAAACTAGGACAAATTACAGGAGCAGTACG
TCTCGACTATCAAGAACGCTTATCATGGAGACACTTGGAAACTAGCTAAACCTTCAATCTTCTGAGCGATTGAG
CTTATCGTCAAGAAAACGACAGGAAATGCCCTCGAATTCTGCTGATATCAGGCCATTACACCATGGAGATTGAGAATT
CCCACAAGTCTGAGTCTTATCAGGAGCGCTCCATGAGTCTGAGTACCTGAGGATACCAACCATATTCAAGAA
CGGATGCTGAAATTGGTCTAATGCCACAATCGCTTATGGGGAGATATCGAAGCAATCCATCTATCGTTCCGTCAGGC
AGACCCGAGATTCTCAATGAGAAATCCACAGCTATGCCCAAAGAACGGCAGGCTCATTATCTCAAGGAAAAT
TCCGAGTACTGAGCTGAGTCTGAGCAGGAAACCATGAGTCTGAGTACCTGAGGAGGAGGAGCTGGGAAATCAACT
ATGATAACAAGCACCAGCTGTTGCAATTACAAACTGACTCCCAATCCAGACAAACAGGAGCAGTCTCTACGAC
AAGGAGCAGTACAGGTGAGGAAGAAGAGTCAACAGAAACGAAACTAACAGGGAAATGGCTTAGTTATCAAGGAGAT
TCTGAAACTTCATCAAGAAAAGGTGTTGCTTAAAGGAAATTGCCCTCTGACCTCCAGGCCAGTCGTAATGACCAAGATT
TCTCGCCCTGCTGAGTACGGAAATTCTGTCAAAAGTACCGGAGAGGAAACAAATTCTCAATCCCTAGAAGTGCAG
CATGCTAGACACTCTGCTGTCATTCAACATCCCTGCAAGACTACGCCCTGGTCCCTTATGAAGTCTCAATGTTGGTT
TGATGAGGATGAGCTAGCACCTTGGCTCCCTGAGAAGCAGAGGATAAGTCCACGAAAATCTCTATGAGAAACTGGCTAAT
GCACAAAAAATGGCAAGTAGTCAAAAGGCTTGTATTCAACAGCTGAGTCT
AAAAACTAAAGCAATTATGGATATCTCTGAGTCTTGGCTTGTATGCCAAACCCACTCTCTATGACTTGTATTGAG
ATTACAAAGCAGCTTATTATGACTATGTTGGGCTTGGCAATGGTCTGCTAGGCAGGCCAATCTCTATGCCCTAGC
ACTGCGTCTGATCAATTGAGAAAGAGCAATTCAAGGTTGCTGCTGTTTATCTGAGTATGAGGACCAAGTCTTGAAGCCC
AGCACGATTGGCAAGGGCTGGCGCAGGCCAAAAGATGAGCTAGAGCTCATGACCATCCACAAAGAGTAAAGGGCTGG
AGTTCCCTACGCTCTTATCTCAATTGGATCAAGATTCAACAGGAAACTCTGAGTACCTGAGTACCTGAGTACCT
AATGGCTCTGCTGCAAAATATATTGCAAGATGGAGACAGGGCAGTAGAGAAGACCAACTATCTAAACCATCAAACCTCCA
TCTCTAGTCTGAGCTATAGGAGAAGAGGAAATTACAGCTAGCAAGCTATTCTGAGCAGTGCCTTGTGTTGCT
ATGACGCCGGCTGAGAAAAGCTCTATCTGTCGGCAAGGGTTCTGAGAAGCTGGAATCAAGGAATACCCACGCC
AAAAATGGGAAACTAAATAGCAATACTAGACTGCAAGCAGGAATTCCAAGATGGCTTGGCTATCAAGTAAAGTGT
TTA

CTAAGGACAAGCTCAACTTAGTTATCGTTTATGGCAAGATCAGTGACCAGAGAAGCTACGGAGAGTTGAAACCAA
GAGTCCTCTCCAAGATACTGCCAACGAGACAATCGCAGTCAGATACCATCAAAGAAGCTCTGAAATGCTGAAGGAGGT
GGAGTTTATAACTCTCACCGCGCAGCTATTGAACTTCTAGTGTCAAACCCAAAGTCAAATCAAGAAATTCTACGAAC
CAGTTATGGATATGGAAGGTCAGATTGCTGGTCAAGGTCAGTCAGTAGGCAAGAAAATCAGCTTCGATITGCCAGATT
TTCAACCAAGAAAAGGTAACGGAGCTGAGATTGGTAGTGTACTCAGAACACTCATGAGAGAAATTGACCTCAGCCAGCA
ACTAACCCCTGCTAGCCTAACAGAACACTCAAACAGTCAGAGAACATCTGCTAACCGACCATCTTATCGCAGAGAACCTTCCATG
CTCAACGAGACCAAAAGAGTCAGGAAGACTTTGTTGCGTGTATCTGCTTACGAAACAAATTG
TTCTGTTGACTACAAGACAGCCCTATGATGAACCAAGTCACCTGAGACCCTATGTTGCGTCAAGTTAGCTATACGA
AGAGCTTATCAGGCTATTGAAATTACTTGTATTACTCGTAAAGACGAGGTTCAAGTTGTA
AAAGTATAA

4109.1 (SEQ. ID. NO. 221)

ATGGAACCTGCTCGCCATGCTGAAAGCTTGGGAGTAGATGCTATTGCAACGATTCCACCAATTATTCCGCTTGCCAGAATA
CTCAGTTGCCAAATACTGGAACGATATCAGTTCTGCAGCTCAAACACAGACTACGTGATTACAACATTCTCAATTGGCAG
GGGTGCTTGAACGCTTACACAGAAATGTTGAAAAATCTCGTGTATCGGTGTAGAGAAACTCTTCTATGCCAGTT
CAAGATATCCAACACCTTGTCAAGCCTTGGTGGAGAACCATATCGCTTAAATGGTCTGTATGAGCAGTTCTAGGAGAC
GCCTCATGGGGCTAGGGCTGGTATCGGTGTACTTATGGTCTATGCCAGAACTCTTGAACACTCAATCAGTTGATTGCG
GATAAGGACTAGAACAGCGCTGAATTGAGTATGCTATCACGCAATCATTGAAACTCACTTCTGCTATGGAAATA
TGACGGTCTATCAAAGAAGTCTGAAAATCAATGAAGGCTTGAATTGGATCTGTTGCTTACCAATTGACACCAGTGA
GAAGAAGATCGTCCAGTTGAGAAGCGCTGCTGCTTGTAAACCAAGGAGCGCTTCTAA

4110.2 (SEQ. ID. NO. 222)

ATGTATAAGACAAGTGTACGAGAGAAGTTAGTATTATTTAAAAATTCTCCAAATCCTGATCTACCAATTGCAA
TTATTCTGCCTCTTTGTTGATCTGCAATGACAGGTCAATACAAACATGACTGGACTGGCTGGTATCTATGGCAACCCAGTA
TCTGGAACTCTTCTTACATTCTAACAGGGATTGTCAGCCTTGGCCTATCAATTGGTCCACATCTTGGTCGAGGCAA
AGGAAGAAGTGTGCTGATTGTTACCAATTATTATGGCTTGGCTTGGGCTATCTGGTCTGGGATGGTACTTTCT
TGGCACCATAATCTGAATCATATTGGTTAGAAGCAGCAGTAGCGGAGTAGCGGTTGCTATCTTGGTTTATCTATC
GGGATTATCCCCTTGCTCTTACGCTATTGCTTCTGCTGGATTGCTGGGTTGACCAAACACTGCCATGTACCTCATG
CTTTGTTACTCCCTCTCAATAGCGGATTAACTATCTTGTGTTACGGTCTGGGTTGCTGGGTTGACCAAACACTGCCATGTACCTCATG
GCTGGTTAGGAACATCTGGCCTACTGGCTCTGGTCTGGGATTCTGGTTCTGGTTTATTAACAGGAGAACGCTCAA
CTTACACCTTGGAGAACATCCCTTAATATGGAAAAATTAGGAAGGACTTGTGTTAGGCTGCTTACCTTGGGAAACT
GTCTTCGGAGTGGCTATCTTCTGGTCTGGTATTGCTAAGTTTCTGGCTTGTGTTAGGCTGCTTACCTTGGGAAACT
GCTATGAACTTCAAGTCTATGAGCCTTCTATGAGTATCTCATGGCTATGGCTATTGCTGTTCTATGAAGTGGG
GCCAAGCGATTGATGCGAAACCTATATTGGTCTAGGAAGATGGACTGCCCTATTTGCGGCTTACCTAACCT
CCTTACATTAGGGAAATGTGCCAGTCTTATGTAACGACCCAAATTATCGATTGACAGTGCCTTTAACTTA
TAGCTTTCTTCCAGTTAGCAGAACCTTGGCGGCGCCGCTCAGGGAAATTGGGGGATAAAGGATAACAGTTACCTT
TTACCTGGTTGCTGGTTATTGGGGCTAGCAATCCCTGTGTACGCTATTGTA

4112.2 (SEQ. ID. NO. 223)

ATGAGTACTTTAGCAAAATAGAACGCTCTTGTAGCGGGTGAAGATGGGATCGGGTCCGCCAGTTAGCTGAACCTC
TCTCTGCACCGACGGCATCCGAAAGTTAGGAAATTAGCCAGAAGTATGAAAAGGACCCAGATTCCAGTTGGC
TTTATTGAGAACAGTGTGCTTATAGATGGTGCCTAATTGCAAGGATTTGAGAATACTCTAAGGGCCT
ATCAACCAAGAGCTTGTCTGGGCTGCCCTTGAGACCTTGTCTTACAAACAGCCGATTACGCCAGATAAAATTGA
TGCCATCCGGAGTTAACGAGTGGAGCCTTGGCAAGGTTGACTGAGTCTGACTAAAGGAAGACGGGAAAAGGA
AGATTGGGGCGCCCCAACCTCTATGTGACTACGGATTATTCTCTAGATTACATGGGATAAACCAATTAGAAGAATTACCA
GTGATTGAGCTTGAGATTCAACCCAAAGAACCAATTATTGGTGAAGGATAAGAAGATGAGAATCAATAA

4113.1 (SEQ. ID. NO. 224)

ATGGATACGATGATTAGTAGATTTCGCCATTATTGAAAGCCTTAAAAGTTGAAACGAAATGGTGGATGACAGTAGC
TGCTGTCACTTCAGTCAGTACTTGTACCTTGGCAATATTGCACTCTTATTTCAATACAGCAGAACTAGCTACAG
ATATTGAAAATATGCGTGTAGTAGTTATACGAAAGGATGTGAAGATAATAGTCAGACAATTGAAAAGAAGGTCA
AACTGTACAAATATGACTACCACAGGTATGTTCTGAAAGAACATGTCTACGGTTAAAGTGTACCTTCTCAAGTA
AAGAAGAACAAATGAAAATTACCGAGATAATGGGAGATAACTGGAAAATTCTGAAAGGAGATGCCAATCCTCTATG
ATGCCCTATATTGAGGGAAACACTCCAAATGATGTAACAAATAGCCGAGATGCTAAAGGAAATTGAGGTGTCTG
GGTCAAGATGGCGTCCAATACAGAAAGACTCTCAAGTTAGCTTCTTATCCGTTGGGACTAGGGATTGCTGCTT
TGTAAATTCTGCACTTGTGATTCAAAATACCATTGCTTACCTTATCCCCTGAGTCGCGAAATTCAAATCATGC
GCTTGGTCTGGAGCTAAACAGTTATCTGGGACCGCTTGTGTTAGAAGGAGCCTTATCGTTATTGGGAGCTATCGCA
CCATCTGTTGGCTTATGTTATCAAATTGTTACCAATCTGTCACAAATCGTTGGTAGGGAAAATCTATCCATGATT
AGTCAGATTATTAGTCGTTGATGATTGCCCTACTATTGTGATTGGGTTTCTATTGGTCAATTGGGATCAGGAATATCC
ATGCGCCGATTCTGAAAGATTAG

4117.1 (SEQ. ID. NO. 225)

ATGAAGAAAGTAAGATTATTTTTAGCTCTGCTATTCTTAGCTAGTCCAGAGGGTGCATGGCTAGTGATGGTACTTG
GCAAGGAAAACAGTATCTGAAAGAAGATGGCAGTCAGCAGCAAATGAGTGGTTTGTACTCATTATCAATCTGGTTC
TATATAAAAGCAGATGCTAACTATGCTGAAAATGAATGGCTAAAGCAAGGTGACCGACTATTCTACCTAAATCTGGTGGCT
ATATGCCAAATCAGAATGGTAGAGAACAGAACGGCTTATTATCTGACCAAGATGGAAAAGATGAAAAGAAATGCTT
GGTAGGAACCTCTATGTTGGTCAACAGGTGCAAAGTAATAGAAGACTGGTCTATGATTCTCAATACGATGCTTGGT
TTATATCAAACAGATGGACAGCACGAGAGAAAAGATGCTCAAATTAAAGGGAGGACTATTCTCAAATCCGGTGG
TTATCTACTGACAAGTGTGATTAGCTGAACTGCTAGTGTGCTAGGGCCTAAAGTACAGCAAGGTTGGCTTGGACA
AACAAATCCAATCTGGTTTACATCAAAGAAAATGGAAACTATGCTGATAAGAAGATGGATTTCGAGAATGGTCACTTAA
TTATCTAAAATCCGGTGGCTACATGGCAGCCAATGAATGGATTGGGATAAGGAATCTGGTTTATCTCAAATTGATGGGA
AAATGGCTAAAAGAACGGTCTACGATTCTCATAGTCAGCTGGTACTACTTCACATCCGGTGGTACATGACAGCCAA
TGATGGATTGGGATAAGGAATCTGGTTTATCTCAAATCTGATGGAAAATAGCTGAAAAGAATGGGCTACGATTCT
CATAGTCAGCTGGTACTACTTCACATGGCAGCCAATGAATGGATTGGGATAAGGAATCTGGTTTAA

CCTCAAATCTGATGGGAAATAGCTGAAAAGAATGGGTCTACGATTCTCATAGTCAGCTTGGTACTACTTCAAATCTGGTGGCTACATGGCGAAAAATGAGACAGTAGATGGTTATCAGCTTGGAAAGCGATGTTAAATGGCTTGGAGGAAAACCTACAAATGAAAATCTGCTTACTATCAAGTAGTGCCTTACGCCAATGTTATGATTCAAGATGGTCAAAAGCTTCCCTATAATGCCAAGGTAGTGCCTAGGCTAGATAAGGATAGAAAAAGTGTGACAAGCCTTGGCTATTACTATTCTGGTTTGTCAAGGCTATAGGAAACAGAAGATTACAAAGCCTAGATGCTAGTAAGGACTTTATCCCCTTATTAGAGAGTGATGGCACCCGGTTTATCTAATGTCAGAATGCTAGTATCCCAGTAGCTCTCATCTTCTGATATGGAAGTAGGCAGAAATTATTATCGGCAGATGGCCTGCATTTGATGGTTTAAGCTGAGAATCCCTCCTTCAAAAGATTAAACAGAGGCTACAAACTACAGTGTGAAAGAAATGGATAAGGTATTAGTTGCTAACACATTAACAATAGCCTTGGAGAACAAAGGGCGTACTTTAAGGAAGGCCAAGAACATTACCATATCAATGCTTTATCTCTTGGCCCATAGTGCCTAGAAAGTAACTGGGGAACAGAAGTAAATTGCCAAGATAAGATAATTCTTGGCATTACAGCCTATGACGACCCCTAACCTTCTGTAAGACATTGATGATGTTGAAGGAATTAGGTGCAACCAAGTGGATTAAGGAAAATTATATCGATAGGGGAAGAACATTCTTGGAAACAAGGCTCTGGTATGAATGGGAATATGCTTCAGACCCTATTGGGGCAAAAATTGCTAGTGTGATGAAATCAATGAGAACAGTGGCAAAGATTAG

4119.2 (SEQ. ID. NO. 226)

ATGAAAAAAAGTATTACAAAATATTGGGCATGGGCTTTGTGTCATCCCCCTTGTACAAGCAATTCTCTATGTGCC
GATGTTTCAAGGAGCCTTACAGTTTACCAACTGGACAGGATTGACTTATAACTACAAAATTGTTGCTTAAACAACTTAA
AGCTCCTTCATGGATCCAAAATTCTATGAAATGCGATTGGCTTACCGCAATCATTGCGATTGCCATGGTGGTGGAGATT
GCACTCGGATCTTCATGGCGGTCTTGAAATTCTAAACAGGCCAACCTCTTCCGTCTGGTTCTTCCCAGCT
GTTTATCTGGTTGACAGTGGCTTGTATCTCAAGCAAGTCTCAACTACGGCTTCCAGGATGGAAATGCCCTCATATT
GAATTCTCAAACCAGCTTTAGGGACTAAGTGGGAGCAATCTTGCAGGCTGTCTTGCCTCTTGGCAAGGGTGGC
TATGCCCATCATCATCTTCTAGCTGGTTGCAATCTATTCAACTGAGATTACAGAGGCAGCAAGGATTGATGGTGCAGACTA
GCAAGCAAGTTCTGGAACATTGACATTGCTTACTCTGCTTACCAAGTGTCTCATGGTCTTATCTCACCCCTAAAGGTGGG
CTGACTGCCTTGTACCAAGTCTTGCCTATGACCGGTGGTGGTCAACATGCCACACCTCACTTGGCTCTTGGTTAA
CTATGCTTAAACACCAATTGGTTATGCCATGCCATTGCCATATCTTGTCTTAAATTGTAGTGAATTGATCAT
CCAATTGAGAGTATCTAAGAAATTGAAATTAA

4119.3 (SEQ. ID. NO. 227)

ATGATGAAACAAGATGAAAGAAAAGCCCTGATTGGCAAATACATTCTATTGATTCTAGGATCGGTTCTGATTTAGTGC
GCCGC
TCCCTGCTACCCCTTITAGTCCCTCAACCCACTAAGGATATTGAGATAATTCTTGGCTTCCAACCAACTTCACATGGG
ACAACCTTAGCGCTCTTACGGATGGGAGGTATTGGAGCTTATTGGAACTCTGCGTCATCACTGTCCTGTTTACTGCA
TAATGATCTTATCCTCATGGCGCCTACTCATCGCTCGCAATATGAGTAAAAGAAAAGCCTTACCATCATGTA
TACCC
TTAACCTCGGAATCTCGTACCTTCCAAGTCATCATGATTCCGATTACGGTTATGAGTAAACTCGGTTGGCAATACC
TTGGTTGATCTGCTACTTGACCTATGCGATTCCACAGACCCCTTCTATGTTGGCTATATCAAATCTGATTCCA
GAAAGTCTGGATGAAGCAGCAGAGATCGATGGGGCTAATCAATTACAACCTATTCCGCATCATCTCCCAATGATGAAAC
CGATGCATGCGACAACCATGATCATGCAATTGCCCCCTTGGTCTGGAAATGACTTCATGTTGCCACTCTTGTCTGAAC
CCGGGAT
TCCAAAATGTTGGACTCTGCTTITGTTCAATACAACCTACGCGAGGCAATATTCAACGACTACGGACCAAGGCTTGC
CTTACGGTGGCATTATCAGTATACCATTGCTATCTCTTCCAACGCCATATCATTTCAGGAATGAGCACGGGGCAGTGA
AGTAA

4119.4 (SEQ. ID. NO. 228)

ATGAAAAGTATTCTTCAGAAATGGGGGAGCCTCCGATGCTGTTCTTTCTAGCTATACTGTTATATCCATTCTTGCA
CAAATTGGATGGGCTTGTGGCTCAGTAGGAATGTTCTATTACTATTTCTTTGCACATACAGTCATTATCCCCT
AAATTCTTCGATTGATTTGCAGTTGCTTGTAGTGTCTTGCACTAGCTGCTTTGCCAGTTAGAACATTCCAATTG
TGAAGAAATTAACTATGCTTTCTTACCCAATATGCAGGGTGGCATCAGAACCGGGCAGAAGTGACCTTCTTAATCCT
AATTATTATGGAATTGTTGTTCTGTATTATGATGTTCTATCTGTTAACAGACCAAGTGAAATTGGTTGAAAGTA
TTCTGTGATTGAGGCTTGTAAATCTTGGTTGAACCTTACTCAAATCGAACCTGCCCTTCTGCTTATATCGCTGGA
GCAATTATCTATCTTACGACTATTAAGGAACTGGAGGCCCTTGGCTTAGTATTGGGCTTCCGGATTGGTTGAGTTTC
CTCTTTCTAGTGATTGGGAGTTGCAATGGGTACTTTAGCTTCTATGGAAGAACGCAATTCTATCTGGATGCTGGAT
GGCCTTGTAAAGCAAATTCTTGGGGTGAAGGGCCATTGACCTATATGAACTCTTATCCTCGGATACATGCTCTTATC
ATGAAACATGCCACAGCTTATTGATGACGATTCTGAGTTACGGAATTGTGGGACTATTATTAGTTGTTCTCTGTG
CTCTCTGCTGCTTGTGATGGATATGAGTCAGGAGTCGGGGAAACGTCGATTATCGGCCTTATCTATCTTCTTACAGT
GTTGCTGTGACGGATTGTTGACTGGCTCTCTGATTAGTCAGTCAGGTTATTCTGCTAGTTATGTCAGTATTCCAT
TGGAGCATCGAATGTGGTATCGGACATGACGGATTAA

4120.1 (SEQ. ID. NO. 229)

ATGTCAAAGATGGATGTTAGAAAATCATTCGACCGATGATGAAGTTGTAAATTGCGTGGCATTATAGCTCTAAAGATGGATGTTAGCAATTGGCCATTGACAGTAGTTGGTAGTTGTTCTTGATTATGGGACAATTGCCGTCGAAGGATTAATAAGAGCATTGCTAGTGTGTTGGAGCTAATTGGACAGAGCCGTTATGCAAGTATATTCAAGGAACTTGTGCTATTATGGGCTAATTCTCTGTTTCAATTGCTATTCTATGCTAAGAATAGCGGAGTAGAGGCTTACCGCTGGAGTCTATCTGTATTCGATTCTTATTGCTAAGATCATCTTATATCCCTAACAGGTGAGGCCATTGGGGAGCTATTAGTAAAGTTGGTTGGAGGCCAAGGAATTATCGGTGCTATCATTAGGTTGGTAGTAGGAAGTATTATACCTCTTATAAAGAGAAAATTGTTATTAAAGATGCCAGAACAAAGTCCACAAGCTATTGCCAACAGTTGAAGCAATGATTCCAGCATTGTAATTCTTATCTCTATGATTGTATATTTAGCGAAGTCATTGACTAATGGCGAACATTCTAGAAGAATGATTATCTGCTATTCAAGTCCGTTGCAAGGTTAACTGGACTTTGTATGGGCTATTGGAAATTGCTATTCTTATATCATTGTTGGTGGTTGGTGTCAATGGCAATCGGAGTAGTAAATTGGAGTAGTGCACGCTGCTTTATCTAACCTGATGCTAAATAAGCTATGTTAGCCTCTGCTAATCTATCATTAAAGATCGTCACATATTGTTACTCAACAATTAGATTCTATTCTATCAGGTTAGGGATTACGTTGGCTTGTAGTTGCCATGCTTTGCAGCAAATCAAAACAATACCAAGCCTAGGAAAAGTTGCAGCTTCCAGCAATATTTAACGTAATGAGCCAGTTGATTTGGATTCCGATTGCTATGAATCCGTTATGTTGTCACCTTCTATTCTGTTCCGACTTGCAGCTGTGATAGTATATGGAGCTATTGCAACAGGTTCTAGCAGGCCATTCTCAGGGTAACATTGCCTTGGAGTACACCAGCTATTATCAGGATTGGTGGGGATGCGAAGGAGTATTACTCAGTGGTGTATATTAGCGATGCTACATTGGTTTATTCTCATTCTTAAAGTACAGGATCGTTAGCTTACAAAATGAAATCAAACAACTTGTAG

4121.2 (SEQ. ID. NO. 230)

ATGAAGAAAAAGGACTTAGTAGACCAACTAGTCTCAGAGATCGAGACGGGAAAGTCAGGACACTGGAAATACGGTCAT
GGAGCTCAGGTAATCACACCTTGCACAGGAATTGTACCAAGCTTACTACAGTAAATTGCTAGAGACAGATC
CTTATATCACCTCAGGACGCCATCTGTAGTACCCAAGGACGCCAATCAAAGGTGACAGCCAGTGCAGTGGCAG
TGAAGTGGAGAGTTGCAGAGAGATATCCTGTTGCAGGGGGATGGATGTCAGTGA

4122.1 (SEQ. ID. NO. 231)

ATGAAGAAAAGATACTAGTCTGACAGCTTGCTAGCCTTGTAGCTAGCAGCTTACAAGAAAAACAAAAATGAAG
ATGGAGAAAACAAGACAGAACAGCAGCCAAAGCTGATCGAACAGCTCGGTAGTAAGTCTCAAGGAGCTGCCAGAAGAAA
GCAGAAGTGGTCAATAAAGGTGATTACTACAGCATTCAAGGGAAATACGTAATCGTACGCCAAACAAACACTATCCA
TTGTCTAAAGACTATAATCAGGGAAAATCCAACAGCCAAGGCAGAGTTGGTCAAACTCATCAAAAGCGATGCAAGAGGA
GGTTCCCTATTAGTGTACATTACAGTGGTTAGAAGTTAGAAACTCAGACCAAGCTATCAAGATTATGTCACCAAGA
TGAAAGGCAGCAGCTGACCGTTACTCTGCCGTCTGGCTATAGCGAACACCAGACAGGCTGGCCTTGTGATTGG
ACTGATGGTGTGTTGGTACAGAAGAAAAGCAGCCAATGGCTTGTGATCATGCAAGCTGATTATGGCTTGTGTCGTTA
TCTCAAAGGCAAGGAAAGGAAACAGGCTATAATGGCTGAAGAATGGCACCTGCGTTATGTAGGAAAGAAGCTAAAGAAAT
TGCTGCAAGTGGTCTCAGTTGGAAGAATACTATGGTTGAAGGCGGAGACTACGTCGATTAA

4125.6 (SEQ. ID. NO. 232)

ATCGTAAATTCTTAATTATTTGCTACCAAGTTTGACCATTCAAAAGTCGTTAGCACAGAAAAGAAGTCGTCTA
TACTTCGAAAAGAAATTTATTACCTTACAATCTGACTTGGTATTATTTAGAGAAAATTAAGTTCTCCATGGTTATGG
AGAGGTTCCGTGTTATCGGAATGAAGATTTAGTGTGGAATTCTGGAAATTGACTCCAAAACAAGTCTTCAAAACCGAG
TGGCGCTTAAATAAACAGGAATTCCAGTTAAGTCTACATCAAAATCATCAATTATAGCTGCCAGAACAGATTTTATATGA
TCAATCAGAGGTAACTCCAACAATAAAAAGTATGGTTAGAATCTGACTTTAACTGTCACAAATAGTCCTTATGATTAAA
GAAGTGAATCATCCTTATCGCTTATCGCAAGTATCAATCGACAAGACCATGTTGAGAAGGAAGAGAATTCTACATA
TTGATCAGGCTGGATGGTAGCTAAGAATCAACTCTGAAAGAAGATAATCGGATGAGTAAAGTTCAAGGAATGTTATCTGA
AAAATATCAGAAAGATTCTTCTCTTATGGTAAAGCAACTGACTCTGGAAAAGAAGCTGATCATCAAGATGAAAAG
ATGTATGCCAGCGCTTGGAAACTCTTCTCTTATACCGAAGAAAATAATGAGGGCTTTATCAGTTAGATAC
GACTGAAAATACGTATCTGAGTCATGATTTCCAGGTTCTATAAACAGAGGGAACTGGTAGTCTCTTAAAGAAG
GATAATAAGAATATTCTTAAAGGATTAAAGCAGTATCAAAAGAATCTGATAATGTAGCTATAATCTATTGGAT
ATTACATTCAAACCAATCTGATGCCACATTCAAATCAAGATGTCTGCCATTATGGAGATGATTGGATCCAAAAGAAA
ATTGATTCTCTAAGATGGCCGGAGTTATGGAGCTATTATAATCAAATGGATTGTGCTAGAGTCTTGA
CAGATTGATAGTCAGGAAATTGCCAAGGTTGTTCTGTTAAAGTAG
CTCATAAAATGGAGATGCCATTAAGCATGATACGGGTGTTCTATGCAGATTCTCATTATCTTCTATTTC
CTAAGAATTCTGATTATGATACGATTCTAAGATAGCCAAGGATGTTATGAGGTTCTAAATGA

4125.7 (SEQ. ID. NO. 233)

ATGAAAACAAAATAATGGTTAATTAAAAATCCTTCTATGGTTATTATTTATCTTTCTTGTGACAGGATTCCAGTAT
TTCTATTCTGGAATAACTCAGGAGGAAGTCAGCAAATCAACTATACTGAGTTGGTACAAGAAATTACCGATGGAATGTA
AAGAATTAACTACCAACCAATGGTAGTGTATCGAAGTTCTGGTCTATAAAAATCTAAAACAAGTAAAGAAGAAC
AGGTATTCACTTTCACGCCATCTTACTAAGGTAGAGAAAATTACAGCACTATTCTCCTGAGATACTACCGTATCAG
AATTGCAAAACTTGCTACTGACCATAAGCAGAAGTAACGTGAAAGCATGAAAGTTCAAGTGGTATATGGATTAACTACT
CGTATCCATTGCCCCATTGGAATTCTTCTCTTCTATCTGATGGAAATATGGAGGAGGCAATGGCCGTAACTC
CAATGAGTTGGACGCTAGTAAGGCTAAAGCAGCAAAATAAGAAGATATTAAAGAAGATTCTTCAAGATTTCAAGATGTTGCTGGAGCTGA
GGAAGAAAACAAGAACACTGTTGAAGTTGAGTTCTTAAAGGATCACAACAGATTCAAAACTTGGAGGCCGTATTCCA
GCAGGTGTTCTTGGAGGGACCTCCGGGACAGGTAAAACCTTGTGCTAAAGGAGTCGCTGGAGAACAGCAGGTGTC
TCTTGTATCTCAGGTTCTGACTTGTAGAAATGTTGCGAGTTGGAGCTAGTCGTTGCTCTCTTGTGAGGATGCCA
AAAAGCAGCACCAGCTACCTTATCGATGAAATTGATGCTGTTGGACGTCACCGTGGAGTCGGTCTCGGCGGAGGTA
TGACGAACGCTAACAACTTGAACCAACTTGTGATTGAGATGTTGAGGGAAATGAGGGATTATCGTCATCGCT
GGCACAAACCGTTAGATGTTACTGACCCCTGCCCTTGTGCTCCAGGACGTTGTAGAAAAGTATTGGTTGGCTCCTGA
TGTAAAGTCGTAAGCAATTGAAAGTTCACTGCTAAGAATAAGC
CTTAGCAGAAGATGTTGAAATTAGTGGCTCAACAAACTCCAGGTTGTTGGTCTGATTAGAGAATGTCTGAAT
GAAGCAGTTAGTGTGCTGTCGCAATAATCGATAATTGATGCTCAGATATTGATGAAAGCAGAAGATAGAGTTATTG
CTGGACCTCTAAGAAAAGATAAGCAGCTTCAACAAAAGAACGAGAAATTGTTGCTTACCATGAGGAGCACATACCATTTGT
TGGCTAGTCTGCTGATGCTCGGTTGCTCATAAGGTTACAATTGATCACCAGCGGCCGTGAGGGAGATACTGATGTC
TCTCTAAAGGAGTCAATGCTTCTATCTAAAGAAGATGAAAGGAGCTTGGCTTAAGGGAGCTGTTGAGGTTGAGCTGA
AGAAATTATCTTAAATGCTAAACACAGGAGCTCAACAGACTTGTAAACAAGGACACAAATGGCACGTGCAATGGTACA
GAGTACGGTATGAGTAAAAACTGGCCAGTACAATATGAGGAAACCATGCTATGCTTGTGACAGAGTCTCTAAAAT
CAATTTCAGAACAAACGCTTATGAAATTGATGAAAGAGGTTGTTCTTAAATGAGGACAGAAATAAGCTGCTGAAAT
TATTCAGTCAAATCTGAAACTCACAAGTTAATTGCAAGCATTATTGAAATGAGAACATTGGATAGTACACAAATTAAA
GCTCTTACGAAACAGGAAAGATGCCGTGAAAGCAGTAGAAGAGGAATCTCATGCACTATCTTATGATGAAAGTAAAGTCAAAA
ATGAATGAGGAAAATAA

4125.10 (SEQ. ID. NO. 234)

ATGAGGGACCAGATTTTAAATCATTTCTCAAGAAGGGATTTCAAAAGCATGCTAAGGCCGTTAGCTCTTCTGG
TGGATTAGATCCATGTTCTATTAAGGTATTGCTACTTATCAAAAAGAGTTAGAGATTGATTCTAGCTCATGTGA
ATCATAAAGCAGAGAATTGATCAGATTGGAAAGAAAAGGATTAAGGAAGTTGCTGCTGAAGCAGACTCCATTAT
CAGCAATTTCAGGAGATTTCAGAACAGCGCTGCACGAAATTTCGTTATGATTTCAGAGGTCATGAAAAGACAG
GTGCGACAGCTTACTGCCCACCATGCTGATGTCAGGTGAAACGATTGTTATGCGCTGATTGAGGAACCTGCTTGC
CGCTATCTCAGGAATTAGGAGAAGCAAGTAGTCGGAGAGATAGAAATCATTTCGAAATTCGATTGAGGAAATTCTTAC
ACTTCCATTCAATTTCAGGAGAAGATACATCAAATCAGGAGAATCATTTCGAAATTCGATTGAGGAAATTCTTAC
CAGAATTGGAAAAGAAAATCTCGATTAGGGATGCAATTCTAGGCAATTGGCAATGAAATTAGATTGATTGCA
AGCTGAATTATCTAACAAATTAAATGAGGAAAGATTACAGCAGTTATTCTACTGAGTCTACACAAAGAGTTACTC
AAACTTATCTGAATGCTTCCAGATTGAAATCTACAAAGCTCAGTTGCTGAAGTTGAGCAGATTAAAATCTAAAGC
CAGTATGTCATCCGATTAAAATGGCTATGAAATTGATAAAAGAGTACCAACAGTTGAGATTGAAAATCAGTCCGAGG
CTGATGAAAGGAAGATGAACTGTTACACTATCAAATCAGGTTATCAAGGATATTCTTCTGGACTTCCA

TTAGAAGGTGAATTAAATTCAACAAATACCTGTTCACGTGAAACATCCATACACATTGTCATCGAAAAACAGGAGATGTTT
TGATTAaaaATGGGCATAGAAAAAAACTCAGACGTTATTATTGAGCAATTGGTGAATTGCTCAATTGGAAATTGCG
ACCAATAATTGAGTAAAAAACGAAAATGATATAATGAACACTGTACTTATAGAAAAATAGATAGGTAA

4126.1 (SEQ. ID. NO. 235)

ATGAAGCGTCTCTCTTTAGTTAGAATGGTTATTCCATCTTCTGGCTTCTCATTCTCCTAGCTCTGGTGGAACTTTCT
ACTATCAATCAAGTTCTCAGCCATTGAGGCCACCATTGAGGGCAACAGCAAACGACCATCAGCCAGACTAGCCACTTTAT
TCAGTCTTATATCAAAAACCTAGAAACCACCTCGACTGGTTGACCCAGCAGACGGATGTTCTGGCCTATGCTGAGAATCCC
AGTCAAGACAAGGTGAGGGAAATCCGAGATTGTTGACCATCTTGAGTCAGATAAGGACTTGAAAACCTGTTGCTGG
TGACCCAATCTGGTCAAGGTCTTACAGATGACAGTGTGAGATGAAAACCTCTCTGATATGATGGCTGAGGATTGGTA
CCAAAAGGCCATTCATCAGGGAGCTATGCCCTGTTGACTCCAGCTCGTAAATCAGATAGTCAGTGGTCAATTCTGTCAC
AAGAACTTGTGATGCAAGGGAGCAATCTGGTGTGCTTGGATATTCTTATGAAACTCTGGAAAGCCTATCTCAAT
CAACTCCAGTTGGGAGCAGGGCTTGCCTTATTATCAATGAAAACCATGAATTGTCACATCCTCAACACAGTTA
TAGTTCTGCTAGCAAAATGGAGGCTATGAAACCCCTACATCGATACAGGTAGGGTATACTCTCTGGTCAAAATCCTAC
AGTCAAGAGAAGAGTGCAGGAACTGATTGGACGGTGTGGCTGAGGTGACCAAGTCTTGGCAACCTATTCCGCTGGGCT
TCTTGTGGACCTTGTGTTGGCCAGTGTACATCTCTTGTCTGCTGTTAGTGTGGTCAATTCTGAGGATTGGTA
AAACGACCCGTCAGTACCAACTTCAAGCCCTTCAGGCCAGATTAATCCACATTCTCTATAACACTTGGACACCATCATC
TGATGGCTGAATTCTGATAGTCAGCGAGTGGCTGAGGTGACCAAGTCTTGGCAACCTATTCCGCTGGGCTCAATCA
AGGCAAGGACTGTGATTGCTCTGACGAAATCAATCATGTCAGGCCAGTATCTTATCAGGAAACAGCTATGGAGATA
AGCTGGAATAAGAAATTATGAAAATGTTGCTTGTATAATTAGTCTACCCAAGCTGGTCTACAACCCCTGTAGAAAAT
GCTCTTACCATGGCATTAGGAAAAGGAAGGTCAAGGGCCATATTAAACTTCTGTCAGAAACAGGATTGGGATTGGTCA
TCCGTATTGAGGATGATGGCTGGCTCCAAGATGCTGGTATAGTAGTCAAAGTCAACTCAAACGTGGGGAGTTGGTCT
TCAAAATGTCGATCAACGGCTCAAATTCTCATTGGAGCAATTACCATATGAAGATTGATTCTAGACCCAAAAGGGACG
AAAGTGAATATATAATAGAATAGAAACTAGCTAA

4126.7 (SEQ. ID. NO. 236)

ATGAAGCGTCTCTCTTTAGTTAGAATGGTTATTCCATCTTCTGGCTTCTCATTCTCCTAGCTCTGGTGGAACTTTCT
ACTATCAATCAAGTTCTCAGCCATTGAGGCCACCATTGAGGGCAACAGCAAACGACCATCAGCCAGACTAGCCACTTTAT
TCAGTCTTATATCAAAAACCTAGAAACCACCTCGACTGGTTGACCCAGCAGACGGATGTTCTGGCCTATGCTGAGAATCCC
AGTCAAGACAAGGTGAGGGAAATCCGAGATTGTTGACCATCTTGAGTCAGATAAGGACTTGAAAACCTGTTGCTGG
TGACCCAATCTGGTCAAGGTCTTACAGATGACAGTGTGAGATGAAAACCTCTGATATGATGGCTGAGGATTGGTA
CCAAAAGGCCATTCATCAGGGAGCTATGCCCTGTTGACTCCAGCTCGTAAATCAGATAGTCAGTGGTCAATTCTGTCAC
AAGAACTTGTGATGCAAGGGAGCAATCTGGTGTGCTTGGATATTCTTATGAAACTCTGGAAAGCCTATCTCAAT
CAACTCCAGTTGGGAGCAGGGCTTGCCTTATTATCAATGAAAACCATGAATTGTCACATCCTCAACACAGTTA
TAGTTCTGCTAGCAAAATGGAGGCTATGAAACCCCTACATCGATACAGGTAGGGTATACTCTGGTCAAAATCCTAC
AGTCAAGAGAAGATTGCAAGGAACTGATTGGACGGTGTGGCTGTCATATTGAAAAGTTAGACCAGGTTGGACTCAGC
TCTTGTGGACCTTGTGTTGGGAGTGTACATCTCTTGTCTGCTGTTAGTGTGGTCAATTCTGAGGATTGGTA
CTCCCTTGAAGGATTGAGGAAACCATGTTGAGAATTGCTTCTGGTGTCAAATTCTGTCAGGAAAGTGGTGCCTAT
GAACCTGAGGAGAAGTAACCGCAATTAAATGATGTTGAGTCAGATGTTGAGTCAGTGTAGGGTACATTCTGAGGAGAAG
AAACGACCCGTCAGTACCAACTTCAAGCCCTTCAGGCCAGATTAATCCACATTCTCTATAACACTTGGACACCATCATC
TGATGGCTGAATTCTGATAGTCAGCGAGTGGCTGAGGTGACCAAGTCTTGGCAACCTATTCCGCTGGGCTCAATCA
AGGCAAGGACTGTGATTGCTCTGACGAAATCAATCATGTCAGGCCAGTATCTTATCAGGAAACAGCTATGGAGATA
AGCTGGAATAAGAAATTATGAAAATGTTGCTTGTATAATTAGTCTACCCAAGCTGGTCTACAACCCCTGTAGAAAAT
GCTCTTACCATGGCATTAGGAAAAGGAAGGTCAAGGGCCATATTAAACTTCTGTCAGAAACAGGATTGGGATTGGTCA
TCCGTATTGAGGATGATGGCTGGCTCCAAGATGCTGGTATAGTAGTCAAAGTCAACTCAAACGTGGGGAGTTGGTCT
TCAAAATGTCGATCAACGGCTCAAATTCTCATTGGAGCAATTACCATATGAAGATTGATTCTAGACCCAAAAGGGACG
AAAGTGAATATATAATAGAATAGAAACTAGCTAA

4127.4 (SEQ. ID. NO. 237)

ATGTTTTAAATTATAAGAGAAGCTCTAAAGTCAGCAGGTTGATCAAAATTTATTACAATTTCATTTATCGTTTGGCT
TTCTGATCGAAACTAGCATTACAGTCTCTGGTGTGAATGCCAATAGCTGAATGTTAAGTGGATTATCCTCTTAAACAT
GTGAGCTCGTGTGGGAATGCCCTAAAAAAACTTTCTGATTTGCCCTAGGAGITAGTCCCTATATCACCGCTCTATTG
TGTCCAACCTTGCAAAATGGATATTACCCAAGTTGAGAGTGGGAAACAAGGGAAAGTAGGTGAGAATGAAATTGAA
CAAGCTACTCGTATATTGCTAGTTCTGCTTGTGAATCTATCGGATTACAGCTGGTTAAACCTTGGCTGGAGCT
CAATTGATTAAAACCTGTTAACCTCACAAGTTCTGACGATTGGTATCATCTTAAACGCTGGTAGATGATTGTCACCTGG
TTGGGTGAGCAAAATTACAGATAAGGGATACGGAAACGGTGTTCCTGATGTTCTGCGGGATTGTTCTCAATTCCAGA
GATGATTACGGGCATCTATGTTGACTACTTGTGAACTGCTTCAAGTGGCTATCATCTATCATCTTGTAAATCATTT
GATTATTACTGATTGTTGATTATTACTTACAACCTATGTTCAACAAGCAGAAATACAAAATCCAATCCAATATACTAAGG
TTGCAACAGGTGCTCCATCTAGCTCTACCTTCCGTTAAAGTAAACCTGCTGGAGTTATCCCTGTTATCTTGCCTGG
TTACTGCAAGGGCTGCGCTATTCTCTGAGTTGGTACGGTACATGTTGCTGGGTTGGGAGGTTAGCACAAGAGATG
TTGGCAACTACTCTCCAACGGTATTGCTGATGTTGTTGATTATTCTTACATTCTTACGTTGATTCTGAGGATTGGT
ATCTGAAAAAGCAGCAGAGACCTACAAAAGAGTGGTGCCTATATCCATGGAGTTGCTCTGGTAAAGGTACAGAAGATA
TATGTCATAACTCTTCCGTCCTGCAACTGTTGGTCCCTTGGTGTGA

4127.5 (SEQ. ID. NO. 238)

ATGGATATTAGACAAGTACTGAAACCATCGCCATGATTGAGGAGAAAACCTCGATATTAGAACCACTACCATGGGGATT
CTCTTGGACTGTATCGATCCAGATATCAATCGTGTGCGGAGAAAATCTAATCAAAAATACGACAAAGCCGCTAATT
AGTAGCTGTTGGTATGAAATTGCGCTGAGTTGGGAACTCTCTACGTTAAACGCTGTATCGGTGACACCTATTCTCTGA
TTGGGGCAGCGACAGATGCGACGGACTACGTGGTCTGGAAAAGCGCTTGATAAGGCTGCGAAAGAGATGGTGTGGACT
TTATTGGTGTGTTCTGCTTGTAGTACAAAAGGGAGATGAGATTCTCATCAATTCCATTCTCGCGCTTIG

GCTGAGACGGATAAGGTCTGCTCGTCACTCAATACGGCTCAACCAAGTCGGTATTAAATATGACGGCTGTGGCAGATATGG
GACGAATTCAAGGAAACAGCAAATCTTCAGATATGGAGTGCCAAGTTGGTTGATTCGCTAATGCTGTTGAGGACAA
TCCCATITATGGGGGGTGCCTTCATGGTGTGGGAAACAGCAGATGTTATCATCAATGTCGGAGTTCTGCTCTGGTGTGTA
AACGTGCTTGGAAAAGTCTGGACAGACAGCTTGTAGTAGCCGAAACAGTTAAGAAAAACTGCCTTAAATCACTCG
TATCGCTCAATTGGTTGGTCAATGGCAGTGGAGAGACTGGGTTGGAGTTGGTATTGGTACTGAGATTGGCACCACCAACC
CCTGGGGTGGAGACTGTGGCACGTGTCCTGGAGGAAATGGGCTAGAAACAGTTGGCACGCATGGAACGACGGCTGCCT
TGGCCCTTGTAAACGACCAAGTTAAAAGGGTGGAGTGTGGCTGCAACCAAGTCGGTGTATCTGGTGCCTTATCCCT
GTTTCTGAGGATGAAGGAATGATTGCTGCACTGGAAATGGCTCTTAAATTAGAAAACAGTAAAGCTATGACGGCTATCT
GTTCTGGGATGGATGATTGCTGCACTCCCAGAAGATACGCCCTGTAACACTATTGCGGATGATTGCGGATGAAGCAGC
AATCGGTATTCAACATGAAAACAACAGCTGTTGATCATTCCAAAGGAAAAGAAGGCATATGATTGAGTTGGTGTG
CTATTAGGAACCTGCACCCGTTATGAAGGTTATGGGGCTTGTGCACTTCATCTCGCCGGTGGACAAATCCCAGCACC
AATTCAATAGTTTAAATAA

4128.1 (SEQ. ID. NO. 239)

ATGACACAGATTATTGATGGAAAGCTTACGGGCCAATTGCAAGGGCAGTGGCTAAAAGACTGCAAATTAAAGGA
GAAACAGGTCTAGTGCTGGTTGGTAGTGTATTGGTGGGACAATCCAGCCAGCCAAGTCTACGTTGCAACAAGGAGA
GGTCAGGCCCTTGCGGCTGGTTCCAGTCAGGAAAGTACTACGGGTTCCAGGACGACCTACTAACAGGAAATTGTTAGACCTGTAT
TGCTAAATACAATCAGGATCCAGCTTGGCATGGGATTGGTTCAGTTGCCCCATTACCAAAACACATTGATGAAGAGGGCGTT
CTATTGGCTATTGACCCAGAAAAGGATGTGGATGGTTCCATCCTAAACATGGGGCCTTTGGTCTGGTCTGGTCTACCGTCAT
GATTCTTCGACACCGGCAGGAATTATGAAATGTCCATGAATATGGGATTGACTGGAGGTAaaaATGCACTCGTCATC
GGTCGATCCAATATTGTCGAAACCTATGGCCCAGCTTCTTGCAAAAGAATGCAACAGTAACCTTGAECTCACTCACGTAC
TCATAATCTTCCAAGGTGGCTGCAAAAGCAGATATTCTGGTTTGCAATCGCTGCGGCCAAGTTGTGACTGTCAGTTG
TCAAAACCAGGTGGCTAGTCATTGACGGTGGGATGAAACGGCAGTAAAGGTAAGCTGTCGGGGATGTTGATTATGAGGC
GGTTGCCCCACTTGCTAGCCACATTAGCCAGTCCTGGAGGTGCGGTCTATGACCATTACTATGCTGATGGAGCAAACCT
ATCAGGAGCACTTAGGACATTGGATAGAAAATAA

4128.2 (SEQ. ID. NO. 240)

ATGTCTAAATTAAATCGTATTCATTGGTGGACTGGATTCTGTAGGAATCGGTCAGCACAGATGCTAATAACTTGTCAA
TGCAGGGGTTCCAGATGGAGCTCTGACACACTGGGACACATTCAAAAACAGTTGGTTGAATGTCCTAACATGGCTAA
ATAGGTCTTGGAAATTCTCTCTGGAAACTCCTCTTAAGACTGTAGCAGCTGAAGAACATCCAACACTGGATATGCAACAAAAT
TAGAGGAAGTATCTCTGGTAAGGATATACTGACTGGACACTGGGAAATCATGGACTCAACATTACTGAGCCTTGCATAC
TTCTGGAAACGGATTCCCAGAAGAAATCTGACAAAAATCGAAGAATTCTCAGGACGCAAGGTTATTGTGAAGCCAACAAA
CCTTATTCAAGGAACGGCTGTATCTATGATTGGACCACGTAGATGGAACACTGGAGAGTTGATTATCTACTTCAGCTGA
CCCTGTITTCAGATGCTGCCAACGAAGACATTATTCTTGGATGAATTGTACCGTATCTGTGAATACGCTCGTTGATTAC
CCTTGAGCGCTCTGCCCTTCTGGCCTCATTTGGCTGCCCTTATGTAGGTGAACCCAGGTAACTTCACCTCGTACGCCAAC
GTCTGACTTGGCTGTATCTTCAATTTCACACTGTTTGGATAAATTGATGAGGCTGGATCGATACTTATGCTGTGGGTA
AAATCAACGATATCTTAAACGGTGTGGTATCAACCATGACATGGTCACAACAAAGTCAAATAGTCATGGAATTGATACACT
ATTGAAGACTATGGGACTTGTGAGTTGAAAAGGATTCTCATTCAAAACCTAGTTGACTTGTGATGCCCTTACGGCCATC
GTCGTAATGCTACGGTTACCGTGTGATTGCTGATGAGTTGATGAACGCTAACCTGAAATTATCGCAGCTATGAGAGAGAAT
GACCTTCTTGTGATTACTGCGGACATGGAAATGACCCAAACGTATGCGGAAACGGATCACACTCGGGAAATATTCCATTGTT
GGCCTATAGGCCCTGCCCTTAAAGGAAATGGTCTCATTCAGTAGGACATTTCAGGATATTGAGCAGTGTGCGCATAACT
TTGGTGTGGAAACTGCTATGATTGGGAAAGTTCTAGATAAATTGGTATAA

4129.2 (SEQ. ID. NO. 241)

ATGTTTATTCCATCAGTGTGGAACTGTGACATTTTACTAACTTTAGTAGAAATTCCGGCTTATCCAATTATAAGAAAG
GCGAAATTACAGGCCAGCAGATGCATGAGGATGTCAAACAGCATCAGGCAAAAGCTGGACTCTACAATGGGAGGTTG
GTTTCTTGTGATTACTCTGTGTTGGTGTCTTCTTCGCCATTATTAGTAGCCAATTAGCAATAATGTGGGAATGATTGTTG
TCATCTTGGTCTTGTATGGCTGGCATTTTAGATGACTTCTCAAGGTCTTCGAAATCATGAGGGCTTAATCCTA
AGCAAAAATTAGCTCTCAGCTTCAAGGTGAGATTATCTCTATCTTCTATGAGCGGGTGGCATACTCTGTCTGTTCTT
GTTATCCAGTTCACTGGGATTCTATATTCTCGCTTCTGGCTAGTCGGTTCTAACAGCAGTAACATTGACAGA
CGGTGTTAGCGTTAGCTAGTATTCGGTTGTGATTAGTTGTCGCTATGGAGTTATGCCATGTCAAGGTAGATGG
ATATTCTTAGTGATTCTGCCATGATTGGGGTTGCTCGGTTCTCATCTTAAACCATAAGCCTGCCAAGGTCTTATGG
GTGATGGGAAGTTGGCCATTGGGGAGTGTGGCAGCTATCTCATGGCTCTCCACCAAGAATGGACTCTCTGATTATC
GGAATTGTGATGTTTGAACAACTCTGTATGATGCAACTGATCTTCAACTGACAGGTGAAACGTATTTCCG
TATGACGCCGTGACATCACCAATTGAGCTGGGGATTGCTGGTAAAGGAATCTTGTGGAGCTGGAGTGAAGGTTGACTCT
CTTTTGGGAGTGGGACTCTAGCAAGTCCTGACCCCTAGCAATTATATTGATGTA

4133.1 (SEQ. ID. NO. 242)

4135.2 (SEQ. ID. NO. 243)

ATGCAAACCAAGAAAACACTCGAAGCAGCCGTTGGCTTGAGCACTTAGCCATGACTCAGGATCTATCCTGG
TCCCCATCATGATTGCAAGCAGCCCTGGCTATTCAGCTGAGCAGTTGCACCTACGATATCTCATGTGCGG
GTGGCAACCTTCCAACTCAACAAATACTTTGGGATTTGGACTCCAGTCTGGAGTTGCATCCAGTGG
CGCTCCCTGATTGATTGGCAAAGCCATGGTAGTGGCGCTATGTTGGTGCCTTATCGCATCTGGGATTACGTGG
TTGTTTCAAGGCATCTCTCAAAGTAGCCAATCTCTCCCATCTCGAACAGGATCTGTTTACACCAGATTGGTTAACCT
TGATCCCCTGCGTATTGGAAATATGGAAATAACGTTCCAGAGCCAACCTGGTCAAAGTCTCTGCTTGAGCTATTACTGTT
CTGATTATCCTCTTGTACACATCTTACCAAAGGATTATCAAGTCTATCTTATTTGATTGGTCTGGTTGGAAACTGCC
ATTGCTGCTACTATGGGTTGGGACTCTCTCTCTGTGAGCTCCAGTGTCCATGTCACACTCCACTCTACTTGGG
ATGCCAACCTTGAATTCATCTTATGTGATGTGATATCGCAACGGTCTATGTTGAGTCACACTGGTTGGCG
GCCTGCTGATATCACAAAGGATCCAATCGACAGCACGCCCTCGAACGGATACCGCGCAGAAGGTTGGCGTACTTC
TCGGAGGAATCTTAAACACCTTCCCTAACCCGATTTCACAAAACGTTGGTTGGTAAATTGTCAGGCATCAAAACAGC
CTGCCAATCTACTACGCAGCTGGTTCCCTGGTCTCTGGACTGCTTCTAACAGTTGGCCTTGGCCAAATCATTCCAAGC
TCCTGCTCGGTGGTGCATGCTGTAATGTTGGTTTGTATCAATTCAAGGGATGCAAATCTGCCCGTGTGACTTGC
AACAAATGAACACAACCTCTTATCGCAGCTGTTCAATCGCTGCAGGTGTCGGTCAACAAACAGTAATCTTGTCAAGCAT
GCCGACAGCCTTCAATGTTCTCAAAACGGAATCGTCAGGCCACTCGTATTGTCCTCAATGCCGTATTAAC
ATAAAAAGAAATAA

4136.2 (SEQ. ID. NO. 244)

ATGAAAGATAGAATAAAAAGAATATTACAAGACAAGGGAAAGGTGACTGTTAATGATTGGCTCAGGCTTGGGAAAGAC
AGTTCAGGAGGATTTCTGAGTTGATTAACCTTGTCTTAATGAAAGAAAGCACCAAAATCGTTTGAGAAGATGGTA
GCTGACATTAGAAAATTAAGAAAAAAACATGAGATTACCTCAAGGGGATTTCATGCCATAAAAATGGCTTGGCTTGG
AGTCTGGAAGGCAGGGAGGACGACTTTGTAGGGAAAAATGATGTCACATGCTATTGATGGTATACCGTCAGGTTAG
TGATTAAAGAAAGTCGCTGCCAATAAGGGAACAGCAGCAGAACGCAAATTATTGATATCTAGAACACAGTTGACAA
CAGTTGCGGGAAATGTTCTGGATCAGGAAAAACCTAAGTATGCTGGTATATCGTCAAAAATCAGAAAATCAGTCA
ACCGATTATGTTAAAGAAACAGCCCTAAATTAGAAGGAACAGAACGAAAGTCTCAAGTCTTATCGATAAAATACCAAGCAG
AAACATGATTCTTGTGCGAGTGTCTCGATGTTAGGGACACTCAACGGATGCGGAAATTGATGTTCTGAGGTCTTGG
ATCAATGGACATTGATTCAGGTTTCCAGAACGGTGTGTTAAGGAAGCAGAAAGTGTGCTGATGCTCCGCTCAAAGGAT
ATGGAAGGTCGTCGGATCTAAGAGATGAAATTACCTTACCATGACGGTGCAGATGCTTCTTATTATGTGACCGAGGGTCTG
ATATCAAGGCTCTGAAAATGGCAATCTGGAGTTGGGGTTCACATGCCAGATGTTCTTATTATGTGACCGAGGGTCTG
CTGACAAGGAAGCCTTAACCGCTGCCACTTGTGTTACGTGACAGACGGAGTGTGCGCAATGCTTCCAGAACGACTATCAA
ATGGCATCTGCTCTCAATCCTTAAAGTGGCCCTGACCCAGTCTGTTATT
GGAGATTGATAAACATGTCGTTGTCACATACATTACAAACAGTTACAGGCTTGTGATGACCTATAGC
GATGTCATGATATCTAGCTGGCATGAAAGAAAAGAGAAAAGAAATATCATAAAATTGATCAAGTATCGAACTCATGCC
AAGCTTCTGAAACTTAAAGAAAACATGCGTGTGAAACGTTGGAGCTCTCAATTGATACCAATGAAGCGAAGATTITAGTGG
ATAAAACAAGGTAAGCCTGTTGATATGTTCTCGGAGCGTGTGATTGCGAGCGATGTTGAGTCTTGTGATGGCT
AAATGAAACAGTTGGCGAACATTTCAGCAAGTGGTTGGCTTTATCTATCGAAATTACAGGAGGCTAACGGCTAACAG
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GACATCATGGCTGCTGTTAGGGAGAACCTTATGCAGATGTTGAGTATGTCAGGAGGACTTCAAAAGGCTAACAG
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TTCTGTTACCGTATGATTGGGATTACGGCGTTCTAAGGAAATAGCAGAGCATTGAAACAGTGAATTCCAGAGATTGGC
ACCCAGTCTTCAACCGTGAACGTCGTTGCCATAGAAGCTGAGCGTAAGTCAAGGCTAACAGGCTAACAG
TTGAGTTATTGATTGTTCTGGAGTTGGCTTGCCTGCGATTGAGTATGTCAGGAGGACTTCAAAAGGCTAACAG
GAATACGTGGGTGAAGAGATGATGTCAGTGTGATCAAGTATTGTCAAATTGGCTCTTGTGCAATTGCAAACACAGTGA
AGGCTTGTATTCACTCAATCTGCTGAAATTGTCAGGAGTGTGATTGACTCTCTGTTGAGAAAATCAGGTA
TCACTTCCGACTGGGTAGCAGATCCGTATCCGTGTTGAAAGAGCGGATAAAATGACTGGAGAGATTGATTGTTCTTCA
CCTAGTGAAGTTGATGTTGATTGAAAAGGCTGAAACAGTCTAGTCG
AGTGGCAGAGGGCGTGAATCAATCGTCGTTGGATAAGAAGGAAGACAAGAGAAAATCAGGACGCTCAAATGATAAGCGT
AACGATTCAACAAAAGACAAGAAGAAAAAGGAAAGAACCTTTTACAAGGAAGTAGCTAAGAAAGGAGCCAAGCATGG
CAAAGGGCAGGGAAAGGTCGTCGACAAAATAA

4137.2 (SEQ. ID. NO. 245)

ATGGGCACAACAGGATTACAATAATTGACTTAATTATCTGATTGTTTACTTGCCTGTTGGTGCAGGTATCTTTC
CTCAAAAGAGATGAAAGGAAGGAAAGACTCTTAAAGGAGATGGTCTGGTCTGGTATGTTACTTGCCTATCCATTGTC
CACAATGCTCAGTCGCTGGATCTTCTGGGACTCTGCTGTTGACTCTTATGCTGAGCTTACCTATCTTGCACGGATAGAC
GATGGTAGTAGCTATTCACTGACAATTGTTTATCTTACCTATCTTGCACGGATAGACATCGATACGCCATATGATTACT
GGATAAACGTTTAACTTAAAGCACTTCGTTATTGACACTTCAATTGAGTATTGATGGTCTGTTATTATGATGGTCT
TATGTAACCTCCATCAGCTGGTTATCAGTATTGACAGGAATTGACATCAATTGAGTATTGATGGTCTGTTATT
TGTGTTCTTACTGTTGCTAAATTCAGGTTATTGAGCAGACTTCAAGGTTGATGTTACTGTTGAGTGGTCT
AGCTTATTGACTGTTGCTAAATTAAAGGTTGGTTGGCAGTAGCAGAACATTAGCAACCGGGAAATTCTTGTG
CAAATGAAAACCTTCTGATCTCAATTGCTTCAACCTCCATCTTAAATTGAGTGGTCTGGTTACGGTTACAATCTTGTCT
CCTATGCTICATCTCAAGATTGGTCAACGTTTACTACAACACAAAATTAAAGAAACTTAAAGATGTTGTCACAAAC
GGTGTGTTGTCACTGCAACTGCAACAGTCTTACTGATTGGTACAGGCTGTTACGTATTCTATCAAGTACAAATGCA
TAGTGCAGCTAGCAATATCCCTCAAGACCAAATCTTATGACTTCTTATGCATACCAAGTACAGTACAGGTTGA
TCTTGGCAGCGATTTATGCGATCTCAACTTCAACAGGTTGAACTCTGTTGCAACTTCACTGGACATTGGTATT
CAAGATGTCATTCTAAAATATGTCAGACAATCTGCTGAGCAGAACATTGCAAACTCTGATCTAGCAGTAGGTTATTCTC
AATTGGTGTGTTCCATTGTCATGGCTACTCAGATATTAAATCTGCATACGAATGGTCAATGTTCATGGACTTGTACTTG
GTCTACTTGGGGTGTATTATTCTGGATTGTTCTAAAGGCAAATAACAAAGGTGCTTATGCAAGGCTGATTGATCA
ACCATCGTCATGGTATTATTAAATACTTCCCTCCAAACAGCTGTTAGCTACTGGGCAATTGATTGATTCATCTG
TCAGTAGTTCAAGGTTATTGATCTGTTCTACTGGAAATAAGTATCTGCACCTAAATACACGATTGATATTAC
AGAAATTAAAGCGGATTCAAGTGGGAAGTTCGTCACTAA

4138.1 (SEQ. ID. NO. 246)

ATGAAATTAGAAAAATATAGCAGCTGGATCAGCTGTTATCGTATCCTTGAGTCTATGTCCTATGCACTAAACCAGCA
TCGTTCGCAGGAAAATAAGGACAATAATCGTGTCTTATGTTGAGTGGCAGCCAGTCAGTCAGAAAGTGAACACTGACA
CCAGACCAGGTTAGGCCAGAAAGGAAGGAAATTGTAATCAAATTACAGATCAGGGCTATGTAACGTCA

CA CGGTGACCACTATCATTACTATAATGGAAAGTCCTTATGATGCCCTTTAGTGAAGAACTCTGATGAAGGATCCAA
CTATCACTTAAGACGCTGATATTGCAATGAAGTCAGGGTGGTATATCATCAAGGTCGATGGAAAATATTATGCTAC
CTGAAAGATCGCAGTCAGTCATAATGTCGACTAAAGATGAAATCAATCGTCAAAACAAGAACATGTCAGGATAAT
GAGAAGGTTAACTCTAATGTTGCTGAGCAAGGTCAGGGACGATATACGACAAATGAGGTTATGCTTTAATCCAGCTG
ATTATTATCGAAGATACGGTAATGCTTATATCGTCTCATGGAGGTCACTACATCACTACATTCCAAAAGCGATTATGCTG
AGTGAATTAGCAGCAGCTAAAGCACATCTGGCTGGAAAAAATATGCAACCGAGTCAGTTAACGCTATTCTCAACAGCTAGTG
ACAATAACACGCAATCTGAGCAAAGGATCAACTAGCAAGCCAGCAAATAATCTGAAAATCTCCAGAGTCTTGAGG
AACTCTATGATTCACCTAGGCCAACGTTACAGTGAATCAGATGGCTGGTCTTGCAGTAAAGATTATCAGTCAGTC
CCAAATGGAGTTCGCTTCGACGCCATTACCTTATCCCTACAGCAAGGTTCTGCCTTAGAAGAAAAGATTG
CAGAATGGTGCCTATCAGTGGAACTGGTTCTACAGTTCTACAAATGCAAAACCTAATGAGTAGTGTCTAGTCAGG
CTTCAAGCAATCTCTTACAGACAAGTAAGGAGCTCTTCAGCATCTGATGGTTATATTAAATCTCAAAAGATATC
GTTGAAGAAACGGTACAGCTTATATTGTAAGACATGGTATCATTTCCATTACATTCAAATCAAATTGGCAAC
CGACTCTCCAAACAATAGTCTAGCAACACCTCTCCATCTTCAATCAATCCAGGAACCTCACATGAGAAACATGAAGA
AGATGGATACGGATTGATGCTAATCGTATTATCGTGAAGATGAATCAGGTTTGTATGAGTCACGGAGACCACATCAT
TATTCTCAAGAAGGACTTGACAGAAAGCAAATTAGGTGCGCAAAACATTAG

4139.1 (SEQ. ID. NO. 247)

ATGAAAAAAAGAGCAATAGTGGCAGTCATTGACTGCTTTGATTGGCTGGATCAGTTGGTCAAATCCTATATCGTCAGC
AGATTCCACTGGGTGAAGTGCCTCTGGATCCCCAATTGCTTAGCTGACCTACCTGCAAATCGAGGTGCAGCCTTCT
ATCTTACAACATCAGCAGCTTATCGCTGTCATTACTCTGGTGTGATAGGTCGCTTGGTATTTACATAAACACAT
GGAGGACTTCTGGATGGTCTTGGGTTGACTCTAAATACGCGGGTGTGTTGGAAACTTATTGACAGGGTCAGTCAGG
GCTTGTGTTGATATGTCACCTGACTTATCAACTTGCATTTCAATGTGGCAGATAGCTATCTGACGGTTGGAGTGA
TTATTATTGATGCAATGCTAAAGAGGAAATAATGAAATTAA

4139.5 (SEQ. ID. NO. 248)

ATGAATACAAATCTTCAAGTGGACTGATCATCGATGAAAACGACCGTTTACTTTGCAAAAGGATGGTCA
AAACATGCTTGTCAAGGAAGAAGCCAACATACAGTAGGGGATACGGTCAAAGGTTTGCATACACGGATATGAAGCA
AAAACCCGCTGACAACCTTAAAGTGAETGCCACTCAGGACCAATTGGTGGGACGTGTCACAGAGGTTGTAAGGAC
TTGGGTGTTGTGGATACAGGCCCTCTGACAAGGAAATCGTTGTGTCACTCGATATTCTCCTGAGCTCAAGGAACCTG
GCCCTAAGAAGGGCAGCAAACCTCACATCCGTTGAAGTGGATAAGAAAGACCGTATCTGGGCTCTGGCTTATCAAGAA
GACTTCCAACGCTTGTGCTGCTCTGCCTACAACAAACATGAGAACCCAAACTGGCAGCCATTGTTACCGTCTCAAGCTG
AGGAATTGTTTACCTACCGAAAATAATGCTTGGTTTATTCTACCTAGCCAGCGTACCGAGGCCACGTTGGGG
AAGTATTAGATGCGCGCTTATTGGTCTCGTGAAGTGGACCCACTCTGAACCTCCCTAAACACGCTCTTGAATG
TTGGAAAACGATGCTAGATGATTGACTTATTGAAAGCAATGGGGTTTATGACCTTAAATGACAAGTCATCTCAGA
CGACATCAAGGCAACCTTGGCATTCTAAAGTCAGTTCAAGAAAGCTTGGTCTTATGAGGCTGGTAAATCAAG
CAGGACCAAGTTGGGACAGAGTTGATTAG

4139.8 (SEQ. ID. NO. 249)

ATGAAAGATGTTAGTCATTGAAAAAGTTCAAAAGCCGTTAAACTGGATTGCTTAGCTTATTGATCTGTA
CTCGGTGTTACCTTTATTAAATAGTCAGACTGCAAACCTCACACAGCTGGAGAGCAGGGTGGAAAGTCGATTGAGCCA
ACGAGAGGGCTATCAATGAAAAGAGAAAACCTCCCAAATGCTGATACAGCTCGGAGGAATACCAAGTTGCTAAAA
ATAATTAGACGTGCAAAACATTTGACCGGAAAGACAGAAAATTCTGACTTATTAAAGAAGGGCGCTGGAAAGAAG
CCTACTATTGCAAGTGGAAAGAGAAAATTGAAATTGATATTGATCAATGACCCGACTCTAGCCCTGGCTTAAAG
GGGGGTGACCGCGAACCGGAAGATTACCAAGCCCTGTATCCCTGAACATAAAAGCACACTTGGAGTTCCGACCC
GGGATTGATCAGATTGCTGGATTAGGGTTATCATCCAAAGTTGTTGGTGTCTATTATTGCTAAACACA
TTGCAAGAAAGATATCAAATCATGGACACAGCTACTTATATCCTGTTCAAAAGTGAATTGCAATATCCTCTTGG
AGTGGAGTGGGATATGTAATGCTGTTATCGGAATCTGCTTCTAGTGGGAAGTCTGATAACTGGTTTG
GACAGTTAGGATATCCCTACCAATTATGCTTAGTGAATCAAGAAGTAACATTGGGAAATAACAGATGTTATTCTCCT
GGCTTGCTCTAGCTTCTAGCTTATCGTCAATTGAGTTGCTACTTGTATGCTTCAAGGAAAATGCT
GTCTCTTCTTCACTCATGGGATGTTGGCTTATTGTTGGTATCCTAACGCTCTCAAGGATTGCAACATCTG
ATTCCTTACTTACTTGCCTCAGTGGAGATTATCTGGAAGATTACCTAACGAGATTGATAATGTCGATCTAAATTGGAG
CATGGGAATGGTCTTACTCCTGCTGATTATCTTGTATTGGAAATTCTATTGAAAGATGGGAAGTTCACAGA
AAAAAGAATTGTTAAATAGATTCTAG

4141.1 (SEQ. ID. NO. 250)

ATGATGAAGTCATATTGGATATTGTTAGTACACCAAGCTATTAGTAGCTTAAATGCAATCTAGGATTAGTTCTCAGAA
GAAGAAATTACCTGATATTAAAGGTGAAATTAGGCTTGTGGTTCTAGTTGATCTGGTGTGAGGAATTGAC
AAAATTCTTAAATCCATTGGTACCATGTTGAGCATGCTTCTCATTTATCGGCTGGTGTGCGGAATATGAAAGCAATTGAG
CTGTAGCTTAAACACATATGGCTCAGCTACTGCAATTGTTGAGGCTATGGTGTGTTCAATATCTTAAATGCTCTTITA
CTCGATTAAATATTTTAAACAGGGCACCACACTCTATATGGCATGTTGATGATTGCGTCTTCTAGTTGCTGGCT
TTACTAGCTGCTCTCATCTTACTAGGAGGATTAGCACTCGTATTATTGAGTATTCCCAGCATITGTCGAAAGAATA
TGGTCAATTAACTGGAAATGACAAGGTAGCTTGGTCAATTGAGTTGCTTCTAGTTGGGATATTGGTTGAGTGGTTTACTGGTAC
CTTATCGGTGACAAATCAAACAGAGGACATTAAATTCCAAAGAGTTGCTTCTAGTGTGAGTACTGTTAGT
TACTTATCCATGGCAGTTATTACATTGAGCTATCTTGCAGGGTCAGAATATGAGAAAAGAAAATCAGTAGTGGTA
CAAGTGGCTAGTTATGCTTCAATTAGCAGGTCAATTGAGCAGGGTATTGTTATTGAGTTGCTTCTAGGTTGCTGGCTTATT
TGGCGAAATTGTCAGCCTTAAAGGTTATTCAGAGCGTCTTGTACTTAACTCAAAACCTGCTTGGATTGCTTCTG
ATACTTATGCACCAATGCAGTTCAATTGGATTATCTCTAGTTGTTGGGTTAGTAAGTATGGAATTATGATTGCTT
CAGGAACGGTTGTTATCTTACCAAGGTGTTGCTCATTCTCTGTGGAGCGACTGCAAGGTGTCATTGGGAAATGCACTGGT
GGTGTGCTGGAGGCACTATTGGAGCATTAACTCAAGGTATTGTTAATGAGTTTCTCCAGTCTTAAATGCCAGTTGGGA
GGACTTGGTTCCAAGGATCAACTTCTCAGATGCAAGATTGGTCTATCAGGAATTATTAGGAATTGTTAAATCAATTGG
CTCACAAAGCAGGATTGAGTTGGCTTGTCTTATTCTAGCAGTTGTTGGAGTATCCTTATTAAAAGCCATCTGCAAC
GGAGGAATAA

4142.3 (SEQ. ID. NO. 251)

ATGATTAAAACATTCTCTGCCCTTCGGTCACTCTTTCTATCCCTATCATAACTTATCTTTCCATCTTCAATCT
TAACATTGGCTATCACCAACCTATCTGGCACAGATTATGCCCTCCCTAGCTACTGCAACTATGGCTGCTATTTAAG
TTCTTATTTTCTATCTTTAACAGAAAAAAATAACAAATACGGTTTACTCTGCATTTGCTTACTATCGCTCATA
TTACTATTATCGGAACAGATAAAACCCCTTCTGCATCAAATAAGACTAAACCTTAAATTAGTAACTTGGAACGTCGC
TAATCAAATAGAACGACAACATATTGAGCGAATTTCAGCCATTGGCTATATCCCCTGAACTAGCTACCA
ATATCAGAGGTGAGCAAGAAACAGAGAATCAAACATTGTTCATCAAGTGGACTTCTATGCCCAACTATGATAATTTC
ACTTCTCCACCTACCAATAGTGAATAGCTCTGTGACTGTGATTGTCAGAAAAGTTATGGTTCTATACAGAACGCTAAAC
TTTCATACAAACACGGTTGGGACAATTGATTACATTGAGAAAACAAATACCGAGATATCATTGGCTTGCATACTGCGC
CTCCTCTGCCAGGTTAATGAAATCTGGAGCAGACTAAACATCATTCTATAATGGCTTCAAATATCCAAGGGCT
ATTATTGAGGTGATTTAATGCAACTATGGCTCATGGAGCAGTGCAAAATAAAGCTCTATAGGGAGCATTAAATGAC
TGCCACCTTGAAGAGGAACTTGGAAATGCCAAAGTCCAAAATTTAATGCAACATAGATCATTTTATTGCTTAA
AACCACTACTATGTTAAAGATTAGACATTGTAAGTTCAAAACTCTGATAGATGTATTTACAGAAATCACATTIA
A

4142.4 (SEQ. ID. NO. 252)

ATGAATCCAATCCAAAGATCTGGGCTTATGTCAGCAGAAAGCGACTGAGAAGTTTATTCTGATTTTATTGGCTT
ATTGGCCGAATTCAGCGTGTGACTCTGATGAAGTCCAACAAAACAGTAGAAAGCAATCTTATAATCACTCAATACA
TCTTTTCTATTAAAGAAGATAGAGAATGGTCAGACATTCAAGTGTGAGACCTAGCATCTGTAAGCAAGATTAAGGGCTGG
AAAATGCTCTCTGAACTTGAGACGGTCGCAAACAACTAAAGACAAGGAAGCAGTGAAGTGGCAGAGCGTGGAGCGTGT
ATGATTTATCAGCTGCAGACAAATAACTTGGTAGCTTAACGGCTCTGAGGATTCATCCAAGGATGTAACCTTACCACTGCG
GCTTCAATCTAAAAGAGGGCAGACCTCAAAAGGGGATTCCAAGAAAATCCTTATCCACGAAGAATTGGCTAAGAAG
AACGGCTTTCGCTTCATGACAAGATTGGCTTGGATGCTGGTCACTGTAATCTGAAAAGGCAACAAAGTAGAGTTTGGAGA
TTATCGGCATCTTTCTGGTAAAAACAGAGAAAATTACAGGCTGTCTCTGACTTCATGAAAGGAAATCAAGTCTTACAGAC
TATGAAAGTAGCCAAACCCCTTGGCAATAGTGAAGCTCAAGTCAGTGCAGCACGGCTCTATGTAGAAAATCCTAAGGAAA
TGGACGGACTCATGAAGCAGGTAGAAAATGGCTTGGAAAATCAAGGCTACCAAGTGCAGGAAAAGGAAAACAAGCTTIG
AACAAATCAAAGACTCAGTGCACATTCCAACCTTCTGACCATCTCCCTTATGGGATGTTAGCAGGAGCTGGAGCC
TTAATTCTGGTTTGTCTCTGTTGAGAGAACGGGCTATGAAAGTGGGATTTCATTCGACTTGGAAAAGGCAAGAGCTC
GATCTTCTACAATTCTGTTAGAGGTAGTTGGTATCTTGGAGCTTCTGCCAGCATTTGTCAGGAAAACCGCAATCAC
AACTTACCTACTCCAAACTCTACTAGCAAGTGGAGATCAGGCAAGGTTACAAGATACACTAGCCAAAGCAAGCAGTTATCA
ACTAGCATCTTATCTTGTGAGAATCTATGTTCTAGTTCTGCTTATGTTCTGTTAGCCCTTGTCTTCTATTCTATT
TAGAAAATCACCGAAAGAAATTATCATCTATTAGTTAA

4142.5 (SEQ. ID. NO. 253)

ATGTTACACAACGCATTGCCATGTACAAGGAAGTTCATACTGATTGTCATCTCCGATTATTCTCCTCATGGCGAGC TTGAGTTGGTCGGCTTGTCAATCAAGGGAGCTACTGCCAACGGCTCTCAGGAGACCTTAAAAAATATCACCAATAGCTTCTC CATGCAAAATCAATCCTCGCGTAACCAAGGAACGCCCTGTTGCTGGGAATATCAAGGGTGAAGACATCAAAAAAAATCAC CGAAAACAAAGGCCATTGAGTCTTATGTCAACAGTCAACGCTATCGGAGATTGACTGGATATGACCTGATTGAAACGCCA GAAACCAAGAAGAATCTCACTGCTGATCGTCCAAGCGTTTGGAGTAGCTTGTGATTACAGGTGTCATGACTCCTCTA AAGAAGACAAGTTGTCCTGGTCTTATAAAACTAGTCAAGGAGAGCACTTAAACCAACGACGACAAGGATAAAATCCTCTT GCACAAGGACTTGGCAGCCAAACACGGCTGAAAGTAGGGACAAGGTTAACTGGACTCTAATATCTACGATGCAGATAA TGAAAAAGGAGCCAAGGAAACAGTTGAAGTGCAACTCAAGGGACTCTTGTATGGTCATAATAAGTCAGCTAACCTACTC ACAAGAACCTTACGAAAACACAGCTTACAGACATTACAGTCTGCAAAACTTATGGGATACACAGAAGACAGCAGGATT TATGGGAGCAGCAACCTTCTTGTAAACAGCAGACAAGAACATTGGATGTTATGAAAGAGTTGAATGGCATCAGTGTATCA ACTGGAAAGACTACACACTCGTCAAGAGCTCTAACTACCCAGCTTGTAGCAATCTATCTCTGGTATGTACAAGATGGCC AACCTCTCTCTGGGTAGCTTGTAGCTTCTCAGTTCTCTCCITGCCCTTGTCTAGCCTTGGATCAACGCCGTCGAAG GAAGTGGGAATTCTCTCTCTATCGGCCCTCAAGCAGGAAGTATCTGGTCA ATTCACTACCGAATCTATCTTGTGATTGTCATCTCTAGTTCTGCTTACTCTCTAGCTAATTACACTGCCGTGCAATTGG AAACACTGTCTGGCCAATGTGACTTCAGGTGTTGCCAACACAGGGTAGTAAGGGGGCTCAAGCTCTAACCTGGTGTGGT CGAGAAGTAGATGGTTAGCAAGACCTTGTGAGCCTAGACATTCCATTAGACATCAACTGACTTTATCATCTTGTCT TGCCTGGTCTAGTGGTCTCGTTATGGCGCTTGTCTCAAGCAATCTCCTTAGAAAACACAAAAGAGCTTGTGGATG GTGAATAA

4144.1 (SEQ. ID. NO. 254)

ATGTCACAGGATAAACAAATGAAAGCTTCTCCCTCTGCAGCGAGTTATCAATATCTCATGATTGTCGGTGGGGTTGG
GAGTTGATTTCTGTATGGGTTATCAGGCTGGATTACAATCCAAGGAAACCCTCTGCCTTATCCAGCAGGAG
GCATCTGGGTCCACCTCTTATCTTACAGATTACAGACTGTCGCCATCATCCAGGGGCTTGACCTCGGTGG
CTGGGTCTTATCTACGGCACATCATCGGGACTATCTACAACTATATCGGATCGTATTGGCTGCCCCATTATCTTATC
TAGTCCCCTATACGGAGCTGCCCTTGCCAGTCAGCACGCCACCTACGACAAGTACATCGACTGGCTAGATAA
GGGCAATCGTTTGACCGCTCTTATTTATGATGATTGGCCATTAGCCCAGCTGACTTCTCTGTATGTCGGTCCCT
GACCAAGATGAGCTCAAGCGTACATGACCATCATCTGACCAAACCCCTTACCCCTCGGGTTTACCTACGGCTGA
CCTATATTATGACTTTCTGGAAATGCTTGA

4144.2 (SEQ. ID. NO. 255)

ATGAGAAAATATGTGGTTGTAATCAAGGAAACCTATCTCGACATTCGACTATGGAGTTCTTCTTATGGTATTGCC
GTTCCCTTTAGAATCTCTGAGGAATTGGCATCTCAAGGTTCTTCTATGGCTAAAATAATAAGTGGCAAGTAGTGA
CAACAGTGCATCTGTAGCAGAAGGACTGAAGAATGTAATGGTAACTTCGACTATAAGACGAGCAAGTGCCAAAG
AAGCAATTAAAGAAGAAAATTAAAGGTTATTGACCATTGATCAAGAAGATAGTGTCTAAAGGCAGTTATCATGGCA
AACATCGCTTAAAGTGAATTAAATTGAGGTTACAGGTACACTAATGAACTGCAAATCAGCTTATCGTCCTAAGTGT
TCCTGTCTCAAGAGCAGGAAACGCTTAGCGCAGACAATTCAATTACAGAAAAGATTGATGAAGCCAAGGAAAATAA
AAGTTTATTCAAACAATTGCGAGGCTTAGGATTCTTCTTATATGATTCTGATTACCTATGCCGGTGAACAGCTCA
GGAAGTGGCCAGTAAAAAGGCACCAAAATTATGGAAGTCGTTTTCTAGCATAAAGGGCAAGTCACTATTCTATGCGCGG
ATGATGGCTCTGTTCTAGTGTATTAAACGCATATTGGGATCATGGTGTAGGGTGTGGCTGGCGGTGTTGCTCTTAAAGAT
TTGCCATTCTGGCTCAGTCTGGATTGGGATCACTTGGGAGATCTCATCTCAATAACCTTGTCTTATTGGTATCAGT
CTTTCATGTCAGTCACTGGTCTGGCAGCCTCTAGGATCTATGGTTCTGTCCTGAGGACTAGGGAAAGCCTTGTGCGCTTGT
ATGATTITGATTATGGGTGGTTTTGGAGTGACAGCTCTAGGTGCACTGGTACAATCTCTTGTGAAGATTGGTCTTAT

ATCCCCTTATTCGACCTTCTTATGCCGTTCGAACGATTAATGACTATGCCGGGGAGCAGAAGCATGGATTCACCTGC
TATTACAGTGATTTGCGGTGGTAGCAACAGGATTTATCGGACGCATGTATGCTAGTCTCGTTCTAACCGATGATTAG
GGATTGGAAAACCTTAAACGTGCCTATCTTATAAATAG

4144.3 (SEQ. ID. NO. 256)

ATGACAGAAACCATTAAATTGATGAAGGCTCATCTCAGTGCAGGTTAAAGAGCAAGAAATTCCCAAGTAGACTTAA
ATGAGATTTGACAGCAGCCCAGATGGCATCATCTTGAAGAATTCCAATCCTACTCTGTATTGTTGAGATITGA
GAAGAAAGATGCCCTGTATGAATTGGTACCTCAAGAAGCCATTGCCAGTCTGTGTTTCCCTCTTGTGAGATITGA
ACCGCAGAAAAGGGAGCCGACTTCATACCGACACCTTCAACCCCAAGGTGAGGAGCTCTTGATGATTAGTTCGTCGA
TGCAGCTTGGACAAAACGCCCTGGAGCTGAAGCTGGGATGGTGTGATTATCGGTTGGTCAACATACA
AGTGAAGAAGTGGCAGAGCTTAAACCTACCTGACTACACCTATTCTGTCTTGGGATGGCACTGGGTGTGCAAATCAA
CATCATGATATGAAACCGAGACTGCCACTAGAGAATGTTGCTTTGAGGAAGAATACCAAGAACAGTCAACTGAGGAATCC
AAGCTTATGACCGTGTTCAGGCTGACTATGCTGGGCGCGACCAAGCTGGAGTCAGCGCCTAGCAGAACAGTTGG
TCAAGCTGAACCAAGCTCAACTAGAAAAATCTTGAACAGAACAGAAATTATTGTAG

4146.1 (SEQ. ID. NO. 257)

ATGTTAAAACCTTATTGCTATTGTTGAACAAATTCAAAACGTTCTACAAACCGTCAATTGCTTCATACATGCAAAACACTT
TAUTGACAAAGCTGAAATTGAACTTGTGAAATCAAGGCCATTCTGCTTCACAAACCCAGCTGACAAGCAAGTACCTGCT
GAAATATTGAAATTGCTGCTAAATCGAAGAGGAGATGGCGTATTATCGGTACTCTGAGTATGATCACTCTATTCCAG
CTGTTTGTAGGCGCTCTGCTTGTCTTATGGTATTACCCACTTTGAACAAACCAATCATGATTACAGGTGCTTCTT
ACGGTAGCCTGGTCTCGTGCACATTGCAACTTCGTAACATTGCAACTGCTCTGAAATCAAGGAAATGTTCTTCCA
GATGAAATTCTGCTCTCACACTCTTCAAGCATTAACCCAGTGGCAGCTGGTGTGACCTTGTGATGTTATCAAGAACATTGGA
TGCCATCTTGTGACTCCGTATCTTGTAAAAATCACAGAACAAATTACGTAATGCACAAGAATTACTCGCAAAGATGCTG
AAGACTTGTACTGGAAAATTGTAA

4146.2 (SEQ. ID. NO. 258)

ATGAATACCTATCAATTAAATAATGGAGTAGAAATTCCAGTATTGGGATTGGAACTTTAACGGTAAGGATGGAGAACAG
CCTATCGTCAGTGTAGAAGCCTTGAAGGCTGTTATGTCATATTGATACGGCGCGATTATCAGAATGAAGAACAGTGT
TGGTCAAGCAATCAAAGATAGCGGAGTCCACGTGAAGAAATGTTGTAACCTACCAAGCTTGAATAGTCAGAACCTAT
GAGCAAACCTCGTCAAGCTTGGAAAAATCTATAGAAAAACTGGGCTTGATTATTGGGATTGTTGATTGATTGATTGATTGCTTGGCGAA
CCCAAACCGCTCAGAGAAAATGACGCATGGAAAATCAGCAACTGGCAAGGAGCAGTGGAAAGACCTCTATCAAGA
AGGAAACCTGGTCTATCGGGCTTAGCAATTCTTCCCACATTGGATGCTCTGTTGAACACTATCGCAACTATGTTCTG
CGGTCAATCAAGTGTGCTTGGGCCAGGTGTGATCAAGATCAAGTGTGCTTACTGTGTGAAAAGGAAATTGATG
AGCTGGGGGCCTTGGACAAGGAGAACTGTTGATAGCAAGCAAGTCCAAGAAATAGCACCAAAATACGGAAAATCGGT
TGCTCAGATAGCCTTGGCTGGAGCTTGGCAGAAGGATTACCACTCCAAATCTGTACAACCTCTGTTATCAAGCTA
ATCTGATTGCTTGGAAATTGAACTGAGTCATGAGGAGAGAACCTTAAAACGATTGCTTCAATCGGTGCTCCAC
AGTTGATGATGTGGATTCTAG

4147.1 (SEQ. ID. NO. 259)

ATGAGGTGCAAAATGCTGATCCAATTGCTATTCAACTAGGACCCCTAGCATTGTTGGTATGCCATTGTTATGTTGACAGG
CTGATTCTGCGGTTATTGACCATGAAAGAACCTAGAAAGAACATACCAAGACGATATTAGTATTAGTCAAGATTTAGGAGAGA
TTTGCCATTGGAATGGGTTGGCCATTACGGTGGTTGAAACTGGGCTTGTCTATATCTTGTGACCGTA
AACTCATCAATACTGGGATTCTAGATATTGCGGCCCTAGCGTTATGATTGCTCAAAGTTGGCGTTGGGTAATTTC
TTAACCAAGAAGCTATGGTCAACAGTGGATAATCTGATTATCTACCTGGTTTACCGTGGCTTATCGTGGACAGATGTATATTGAGGG
GAGCTACCGTCAACCGACTTCCATTAGTGTCTATGGAATCTGTTGGCTTGCCTTGTGATTGATTTAGCAGGAAATG
GAAGAGTCTCAGACGAGGTCATATCAGGCCCTTACTTGATTGGTATGGGTTGGCTATGGTATCCAGGATGCGAA
CAGATAGTCTCATGTTCTCGGCTTGCAGTGTCCCAGTGTCTTATCGGCTCGGTATAATGATCGTTATT
ATCAAATCGAAAGAACGGCCCTACTATATTACAGAGGAGAACCTAA

4147.2 (SEQ. ID. NO. 260)

ATGGGTAAATTATCCTCAATCCTTATTAGAACCGTTCAAGGTGAGCTTGCCTTGTGTTAACAGTGTAAAGGCAAACA
AGTTGCACTCAGGCTCAAGATTCTAGATGTTGAGAGAACGATCCGGAGTATGCCAGAGCTGTGAAAAAAACTG
ACAGAACGTTAAGGAGCAGGCTCACAGATTGTTCTGAAACAAAGAACAGGTTGAGTCAGGTGAAATCACTGTGGACAGT
ATACTGCTCAAACAACTATGCTTCAAGCAGACAGAACATCAAAATCAATTAAATAATCTAAGGAGCAATGGC
AAGAAAAGCCGAAGCTTGTGACTCAGAACAGATTGTGATTGATATAACAGAACAAATA

4147.3 (SEQ. ID. NO. 261)

ATGAAAACCTAAATTGATCTTGGGGCTATGCTCTTCTCCCTCCATCCTCTGACCATTATCTGGCTGGATT
TCTATCCTATGGAGATTAGTGGCTAAACTTAAACGAAATCGAGTCTATCTAAAACCAAGAACATTCAATACAATTTCATATC
TTGATGAAATTATCTGACCAATCCTTACTGAGGTCTTACAGATGCCGTTATCTGCTCTGCTCAGCAGCTGGCTGACCAATTTC
GCACTGGTCAAGAATCTTTCATTGGTTCAAGCTAGTAGCTAGTGTACACTGCCAAGTTCTATGTTCTGCAATTAGGATT
GTGAAAAGGACTTTGCTCTTATCGAAAAGTCTCTGGCTCTAGTAGTCTTACCTGTGATGATTGGACTTGGGGAGT
TTGATGGGTTTGGACCAATTCTTACTCTTCTCATCAAATTCTCTTGTGGGAGATGATACCTGGCTTTTGATCCAGCAA
GGATCTGTTATTGATTTGGCAGAGACCTTCTTCTCATGCCCTCCCTCTTGTGAAACCTTCTGAAATTCTTGGC
TATCTGATCTGAAAGCTGAGGAAGTGA

4149.1 (SEQ. ID. NO. 262)

ATGACTTATCATTAACTGAAGAACATCGATATTATGTAATTGGTGCAGGGACACGGCTGGGTTGAGGCTTCTGGCCCTAG
CCGTATGGGCTGTAAGGTCTGCTTGCAGCCATCAATATTGAAATGCTGGCTTCTGATGCCCTGTAATCCCTATCGTGGTTC
TGCCAGGGATTGCTGCTGAGTCAGTGCATGCCCTCGTGGCGAGATGCCAAACCAATTGACAAGACTACATCCAGATG
AACATGCTAAACACAGGGAGGGCCAGCTGCTGCCCTCGTGGCGAGCAAGAACATTAAGGAGAT
CGCAAGACGGTTGAAAACCAAGAAAATCTGACCCCTCGTCAAACCATGATTGATGAGATTGGTGGAGATGGCAAGGTTG
TCGGTGTGCGTACAGCCACCCATCAAGAACATGCTGTAAGGCTGTTATGTGACCGACAGGGACTGCTCCGTGGGAAAT
TATCATCGGAGACCTCAAGTACTCATCAGTCCTAACCAACAGCTGGCTTCTATTAAACCTAGCTGACAATCTCAAGGAACGG

GTCTCGAAATCGTCGTTCAAGACAGGAACCCTCCACGTGTCAAGGCTTCTTATCAATTACGATGTGACAGAAATTCA
CCAGGAGACCAAGTGCTAATCATTCTCATACACTTCACGTGATGAGGGATTATGTCAAGGACCAAGTACCATGCTGGTGA
CCTATACCAATGCTACCGTACATGAGATTATCCAACACCTCCACCGTGCCTATGTTACAGGTGTTCAAGGGAGT
GGGGCCTCGTTACTGTCGTCGATTGAAGACAAGATTGTCGCTTGGCAGAACAGGACCTGACCAACTTCCTGAGCCA
GAAGGGCGCAACTTGAGGAAGTCTATGTCAAGGACTTCAACCGACTGCTGAGGATGTCCAGCGTGAATTGGTGCATT
CCATCAAAGGTTGGAAAATGCAGAGATGTCGGACAGGTTATGCTATTGA
GTATGATATGGTCTTGCTCATCAGTTGCGTGCACCTTGAAACCAAGAAAATCTCAGGTCTTCACTGCTGGTACAGACAA
ATGGAACATCAGGTACCGAAGAGGAGCAGGCCAAGGGATTATGCGGGTATCAATGCGGCTCTGAAAATCCAAGGCAAGC
CTGAATTGATTGAAAGCGCAGTGTGTTATACGGGGTGTGATGTCGACGACTTGGTAGCCAAGGGAACATTGAACCTTA
CCGTCTTGTGACCGACTGCTGAATACCGTCTCATTCCTGCTCATGACAATGCTGATATGCGCTTACTGAGATGGGACCG
AGATTGGCCTTGTGGACCGATGACCGCTGGCTGTTGAATCAAGAAAATCAATTGATAATGAGATGAAGCGCTAGA
CAGTATCAAACACTCAAGCCAGTCAAGGAACCAATGCCAAGGTTGAGGAGATGGGCTCAAACCTTGACCGATGCA
AGCCAAGGAATTCCCTGCCGTCCAGAAGTTTACCAAGATGTGGTGGCTCATGGACCAGCTGCAGAAGACTTGGAT
GACAAGATTATCGAATTGATTGAAACAGAAATCAAGTATGAGGCTATATTCCAACAGGATGGACCAGGTTGCAAGATGA
AACGCATGGAAAGAAAACGCATTCCGGCAATATCGACTGGGATGACATTGATTCTATCGCAACCAGAAGCCCCTGACAGATT
CAAACCTCATCAATCCAGAAACCATCGCCAAAGCGTATTCCGGAGTAAACCCAGCAGATATTCTATTGATGGT
TATCTGGAAGTAAAATCGTAGTATTCTAAAAACTCTCAAAATCAAATGA

4149.2 (SEQ. ID. NO. 263)

ATGAAAGTATTAGCTTGTGATACGTCCAGCAAGGCTTTCTGGCTATTTAGAGGATAAGCAGGTTCTGCCAGACGAC
GATTAATATTAAGAAAATCACAGTATTACTCTTATGCTGCATCGATTGTTGATGCCAAGTTGGATTGGACACCCAAGG
ATTGGACCGAATCGTGTAGCTGAAGGGCCGGTAGCTATACAGGCTTGCAGATTGCGTAGCAACTGCTAAGACCTTAGC
TCACACCCCTGAAACATCGCAGTTGGGTATGTCGAGTCTTGGCTGGCCATCAACAAGAAGGTTGGTGTGCCCCCT
TGATGGATGCCGTGCAATAATGTTATGCCAGGTTATGAAATGCCAACCTGTGATGCCAGAACGGCACCTATCTTT
GAAGAGGCTGAGAAAAGTCAAGGGTACTAGTCAGGTACCTTGTGGAGAAGTTGGCCCCCTTGTGAGCAGATTCAA
AACACTGCCAAGGACTGATTACAAGAAACATTGCCAATGCACTAATCTAGCTCTTGGCTGGACAAGGAAGCAGA
CTCCTTGATGATTGTCGAAATTACCTCAAACGAGTCGAGGAAACTGGCTCAAGAACCATACCGAGTCTGG
GAGTCTTACATTAACCGCTATGA

4149.3 (SEQ. ID. NO. 264)

ATGATAGAAATCAAGCAATTCAACACAGCTGACCTAGCTCAAGGCATCTACGCTGTTATGGCAGCTGTTACCTAGTCA
GTCCTTGGACTCTGGAGCAATCCAAGCAGATCTGCTTCAAGACCAGACTTGGTATGCAATTGCTTATGATGGGGAGAAGT
GATTGGATTCTAGCTGTGCAAGGAGATCTTGTGAGCAGAAGTCTGCAAATCGCTGTCAAAGGAGCTTACAGGTCAG
GGGATTGCGTCAGCTTGTCAATTGCCAGACAGAACGAAATTCCCTGCAAGCAGTCAAATCAACGAGGCGCA
ACGATTTACAAGAAAGAAAAGTACAGTTATCGCTGAGCGAAAGGCCACTACCATGACCCAGTCGAGGAGGCCATTATC
ATGAAGAGAGAAATAGATGAAGGATAG

4152.2 (SEQ. ID. NO. 265)

ATGACAAAACAAGTCTTATTAGTGGATGATGAAGAACACATTCTGAAATTGCTTACTACCATTAAAGTAAGGAAGGCTTT
CTACTCAATTGGTACAAATGGACCGAAGGCCCTAGCTTGGCAGAACACAGAACCCCTTGATTITATCTGCTTGTATCATG
TTACCAATTAGATGGCATGGAAGTTGTAAAGCGGTGAGAGCCAAGGGCTCAAACACTCAAATTATGATGGTTCTGCA
AAAGTGTGATGATTGATAAGGTTGGCTTGGAAATTAGGGCTGATGACTACCTGACCAAGCCTTTAGCCCTAGAGAATT
GCTGGCGCTGTCAAGGCTGCTCAGCGAACTAAAGGAGAACAGAAGGAGATGATTGAGATAATATGCTGACGATT
TTGGCTATTGGACCTTGAAGTATACCTGAGCGTATGAGTCTACAAGCGAATAAGTTACTGAGTTGACCCAAAA
GAATTGAAAGCGATAAAAATCCCTTTTGAAGTTCAAGTTGAAAGTAACCGCCCAATAA

4154.1 (SEQ. ID. NO. 266)

ATGACTACTTTAAAGATGGTTTATGGGGGGTGTGCTGCTCATCAACTGAAAGTGGATGGCAAGAAGGTTGCA
AGGAATTAGTGTGCTGATGTTATGACTGCTGGTGTGCTATGGAGTAGCTCGTAAATTACTTGGAGTTAGGGTAA
TATTATCCAATCATGAGCGATGATTATCACCGTTAAAGAACATAGACTTTGCTGAAATGGGATTCAAGTG
CTCCGTAACCTCTATTGATGCCAGCTATTCACGGCAAGGCTGTTAGCTGAGGCGAATGAGAACAGGATTACAGTTATG
ATAATCTTTGATGAATGTTAAAGAACATGGTATTGAACTCTGATCATCCTATCTCATTTGAAATGCCATTACATTAGTA
CCGAATATGGGGTTGGAAAATAGGAATTGATTGATTCTTGTGCTTGGAGAAGTGTGTTAAACGTTACAAGAT
AAGGTTAAATATTGGATGACTTCAATGAAATCAATAATCAAGCGAATTATCAGGAAGATTGACCCATTACTAACTCAG
GTATTGTATGAGGAAGGTGATAATAGAGAACATTATGATCAAGCAGCACATTACGAATTAGTTGCTCTGCACGAGC
TGAAAATTGGTGTGAGGATAATCCAGATTCTAACATAGGTTGATGATTGCGATGTTGCAATTATCAGTTACTGCA
ATCCTAAGGATATCTAATGGCAATGAAAGCTATGCAAGCGTATTATTGCTGATGTCATGTTAGGTAATATCCT
GAGCATTTCAAGTATTGGACCGAAGGGTATTGAGTTACTGCCAGGATAAAAGAACATTGAGTTACTGGGG
CTGTAGATTACATTGGTTCACTATATGCTTGTATGACTCTCATCGTAAATAATCCTTATTGATTATCTG
AAACAGAACATTAGTGAAGAACATTATGTTAAGGCTCTGAATGGAGTGGCAATTGATCCAGAACGGTTGCTTATGC
GTTAAATTGGTTACAGACCACTACTTACCACTCTTATTGTAAGAACATTGGGGAGCTATGATCAAGTGTGAGCAG
ATGGTATGTCATGATGATTATAAGAACATTGAGCTTACGGTGCCTTATCGTGAATGAAAAGGGTGTAGTTGAAGATGG
TGGTATTGTTAATGGGTTACTCCATGGGATGATTGATTGGTTCACTGGTACCGGTGAATGCGGAAACGTTATGGCT
TTATTGATGATAAAGGAGTAAATGGGAAGGAAAGTTATAATGTTCCCGAAAAATCTTGGCTGGTATAAGGAAGT
TATTICATCTAACGGTGAATCAGTAGAATAG

4154.2 (SEQ. ID. NO. 267)

ATGGATCAACAAAACGGGTTTCTTCTGAAAACCATGTTATGGGACCAATGGGCAAACCTGCTCAGTTAAAGTAGT
ACGTGCTATCACGGCTGAGGTATGGCTGTACCATTAACGTTAGGATCAATGTTTGGTATTGATGATTCTGCCACA
AGCTTCTCATTTGGCAATTGCGAGATATTCTCTGCTCATTTGATAAATTCACTACATTACATGGTGCACACTA
TGCAGACTATGGGTTCTCTATCTCTTATTGCTCTATCACTGCAATGACAAAATTATGCAAGGAGAAC
TCATATGAATCCCTTAATGGTGCCTTGCCTTGTGCTGACAGTACCGCAATCATTGATGGTGTGAA
TGATGAAGACTGTGACAAGTCTAAAAGAACGGTGCAGTAATTGCAAGATGGATGGCAATGGGAAATGAGTCGACGTTG

GGACAACAGGGATTTCACCGCAATCATTATGGCAATTGTGACTGTTCTTATTATCGTATGTGTAAACATAATTGGTT
ATTAAAATGCCTGAAGCTGTTCCAGAAGGAGTTCTCGTGGATTACCGCTTGGTCCGGGATTGTTGTCATTGTTGTT
ATCTTTATCACGGCTCTTCTGTAGCAATGGAACAGATTTAAAGTCATTCGAATTCTCATTTGGTTGTTGATCCAATCTG
ACTAATTCTGGATTGTTTAATGATTATTCTATTGACTCAACTACTTGGATTGAGGTATCCACCGTGCAGAATTGTT
TTGCAATTGTTAGTCAATTGCTCTTGCTAACATGGCTGAAAATGCTGCTGGGGGACTCCGCTGTTGCAAGGTGAAATTCT
AATATGTTGTAATTGCAAGGTGTTCTGGTCAACTTAGGACTATGTTATATATTGCTTTGCCTAAATCTGAACAGCTT
AAAGCAATAGGACGAGCATCTGAGTTCAGCCTTATTAAATATTAAATGAAACCATTAAATTGGATTACCTATTATCTATAA
TCCAGCCTTGGCTATACCAATTATTTAGCACCACATGGTACTGCTACTATTATACGTCAGCAATTCTCTAAACATTATAA
GCCAATTATCGCACAGGTTCCATGGCCAACTCCAGTGGGATTGGAGCTTCTAGGGACAGCAGATTCGAGCTGTATTAA
GTTGCTCTAGTATGTCATTGCACTTGCAGCATTCTAGTCTATCTTCATTATCCGTTGATATGATCAAATTGGTGAAGAAGA
GCAAGGTATCTAA

4155.1 (SEQ. ID. NO. 268)

ATGAAAAAAATTATGTAGTCAGTCAATTCTATTCTAGTAGGATTGGCTTGAGCTTATCCACTTTCAATTATTTT
GTTAATAATAATCTGTTGACGGTTTAATTGTTCTTTGAGGAGGCTATGTTTTTATTTAAGAACAGAGTCAT
TATACAAGGAGTGTAGAACAGATACAGTATGAAACCACCAAGCGGAAGAACAGTTGACAGCTTATGGAACAGATG
CCTGTAGGTGTTATGAAATTGATATTCTTGGAGAGGTTGAGTGGTTAATCCTCATGCTAATTGATTTGACCAAGGA
AGATGGTGTGATTGATTTAGAACAGTGTCAACAGATTACAGGCTTCAAGGAAATCCGCTACTTATGCCAACGTTG
AGAACGGCTTATGCTGTTCATGGATGCTTCTCCGGTGTGTTGATTTGAGATGATCCAGGGAACAGCATAACAGAT
GAATTGGTAACAAGTAGACCAGTGATGGGATTGTCCTGTTGATAATTATGATGATTTGAGGATGAAACTTCTGAGTCAG
ATATTAGTCAAATCAATAGTTTGAGCTAATTGATATCAGAGTTTCAGAAAACACATGATGTTTCTGTCGGGTAAGT
ATGGATCGATTATCTATTACTGACTACACGGTGTGAGGGCTGATGAATGATAAAATTCTGTTATTGATGCTTCTGAG
GAAGAGTCGAAACAGAGACAGTGGCCCTGACCTTAAGTGGGATTCTTATGGCAGTGGAAAATCATGATGAGATAGGA
AAGTGTCTTGTCAATTGAACTTGGCTGAAGTACGGTGGCGACCGAGGTGTTAAGGAAAACGACGAAACGAAAAA
TCCAGTTATTGGTGGGGCTGCTGCTCAATTCAAGCGTACACGGACTCGTACCGCCCTATGATGACAGCTATTICAG
ATAAGATTGGAGTGTAGATCAGGTTTGAGTCGGTCAACAAAATTAGACATGGATGCTTGGGCTCTGCTGAGGTATG
CAGTTGTCGCCAGCAATGTGATTGAAAATAGCTATGCTTATGATGAAGAACAAATGTCAGATATTGAACGAGCTG
TTCAATTCAAGAAAAGAAGGAGTTCAGAAGTTGTTGCTGTTAAGGATGCAATGGGGATGGTGCACCAATCGTTCTTGTG
ATTCTGTAGACCATCAAAGACAGCCTTAACATTATCAAAGAAATTATGATTTATACCCAAACCATGTTATTGACCA
CCATAGAAGGGATCAGGATTTCAGATAATGCGGTTACTTATATCGAAAGTGGTGCAGTACTGCGCAGTGAATTGGTA
ACGGAATTGATTCACTTCAGAATTCTAAGAAAATCGTTGAGTCGTATGCAAGCAAGTGTCTGATGGCTGGTATGATGTT
GGATACTAAAATTCACTCGCAGTAACTAGTCGGACATTGATGTTGCTAGCTATCTAGAACCGCCGGAAAGTGTAGT
ATTGCTATCCAGGAATCGCTGGCAGACATTGAGAAGAATATCGTAGGCTAATGAACTTATTACAGGGCGTAAATTAG
GTTCACTGAGTACTAATAGCAGAGGCTAAGGACATGAAATGCTATGATACAGTTGTTATTAGTAAAGGCAGCAGTCCTCATGTT
AGCCATGTCAGGTTATGAGCAGGATTGTTCTGCGAAGAATACACAAAGGATTATCTCATCTCAGCTGAGTCAGTGA
AACTGAATGTACAACGGATTATGAAAGAGTTAGGCGGGGAGGCCACTTAAATTGGCACAGCTCAATTAAAGGTGAAAC
CTTGTCAAGCAGGGTAAAAACTGACAGAAATTGATTAATGAAATGAAGGAAAAGGAGAAAGAAGAATGA

4156.1 (SEQ. ID. NO. 269)

ATGAAAGAGAAAAATATGTGAAAGAATTGTAATCGTCAGGGCTGGATTTGGCTTTACTTGCGCTCTTATATCA
GGTCTCCCTAGTGGTTACCTCTATTGACTTAAAAGAAGTAGGCCCTGCTACAGTCAGGCCTGATAAGTGTGGCCTTCAA
TTGTGGTCTGGCTCTATTGAGCTCGTAAAACCAGTACTAGTAGTTTAACTTCTTCTTATAGCTAACAGATT
TGGCACGGTGGGCTTGGAGTATCTAGTTATGTGGCTCAAATATACTGGTTCCTTATTGCAACTGTCAAATGAGACG
ACAACAGCTAACCGAGTCTCAGATAATGATATGGTCAAAATAGTCTGGTATTTCCAGTTCTCTCTAGCCTTGGCT
CCGATTGTGAGGAATCTTGTGCTGGGATTGTCCTTAAAGATTTCCGAGGAAGGAGAACTTGGGATTGTAGTCG
GTACGATTGTGTTGCTTATTGCATCAACCAAGTAATTACCTTCTTATTGATTATGGAGGTATGTCGACAGTTCTATCTT
GGACAGCCTAACAGACCCAACCTTGGAAATGTCGATCTGCTTACATGATTGTTAATGGGATTGCTTCTGTTGGCT
CTTGTGGTATTATGAGTCGGACATTAGGAATTCTGTTAAATGAAAGAGAAAAATATGTTGAAAGAATTGTAATCTG
CAGGGCTGATTGGCTTTCATTCTGGCTCTTATTACAGGTTCCCCTAGTGGTACCTTATTTGACTTAAAGAAG
TAGCCCTGCTACAGTCAGGGCTGATAGTGTGCTGGCTTCAATTGTGGTCTGGCTCTATTATTATGGAGCTGTAACACC
AAGTTAGCTAGTTTAATTCTTTTAAAGCTAAAGATTGGCACGTTGGGCTGAGTTATCTAGTTATTGTCGGGTCA
AAATATCTGGTCCATTATTGCAACTGTCAAATGAGACGACAACAGTCAACCAGTCTCAGATAATGATATGGTTCAAA
TAGTTCGGTATTCTCAGTTCTCTGCTAGCCTGCTGGCTCCAGTTGTGAGGAAATCTGTGTCCTGGGATTGTCCTAA
AAAGATTCTGGAGGCAAGGAGAACCTGGATTGTAGTCGGTACAGTTGTGCTTATTGCAACCAAGTAATTAC
CTTCTTATTGATTATGGAGGTATGTCGACAGTTCTATCTGGACAGCCTAACAGACCCAACGTTGGAAATGTCGATCTG
CTTCACATGATTGTAATGGGATGCTTCTGTTGTCCTGGTATTGAGTCGGACATTAGGAATTCTGTGTTAA

4156.4 (SEQ. ID. NO. 270)

ATGGATACACAAAAGATTGAAGCGGCTGAAAAATGATTATCGAGGCTGTAGGAGAGGCAGCTAATCGAGGGCTTGCAG
GAAACACCTGCTGTAGCCCGTATGTATCAAGAGATTTTCAGGTCTTGGTCAAACAGCAGAGGAACATTGTCAAAAT
CCTTGTGAAATTATTGACGATAATATGGTGTAGAAAAGGATATCTTTTCCATACCAGTGTGAACACCCTTGTGCCATT
ATGGTAGAGCGCACATTGCCATCATTCCAGATGGTGTGGCAGGCTTGTCAAGCTAGGCCGTACGGTTGAAGTTATTG
AAAAAACACAAAATTCAAGACGTTGAATATCGAAGTGGCGATGCCATGACTATCTAGGTGCTAAAGGAGCCTTGT
TGTCTATTGAGGCGGAACATATGTGTATGAGTATGCCGTGGTGTAGAAAACCAGGCACTGCAACCTTGACGACAGTAGCTCG
TGCTCATTTGAAACAGATAAGGATCTCCGTGACCAAGCTTATCGTTAATGGGGCTATAA

4157.2 (SEQ. ID. NO. 271)

ATGAAAGACTTGTCTTAAAGAGAAAAGCAGGCCCTTCGTAAGGACTGTCTGGTTATCTGGCTATGTGCTCAATGACCACTT
TGTCTTGTCTGCTTGTCTGGCTTCTAGCCTACCACTAGCTAACACTTACAACATTCTGAAAATCATGGCC
TATCCTTGTGTTGAGAATTACGCTCTGTTACTTTACTTGGGGAGGAAGTGCACCTATATGGAGGCTCCAGACAAGCT
CTTCTCTAGTGGAGAAGAGGAATTAAAGCTCATCTCAAGCGTCAAAGTGCATTCCCTAGTCTTGGCTCTTGAC
GACCTTCTTGTCTTATGGCCCTTATTTAGCAATGGGTTATGGCTTGCAGTTTCTGCTATGTGCTTTATTG
GGGGTAGGAAAATATTCCACTTGTCTAAAGGGCAGCAATTCTACTGAAACTGGACTGGAGGACTATGTTATTC
TCAAGAAAGCAAGCGTAAGCAAGTCTGCTTGTCTTACGAGGTCAAGGGAAATTCAACAGCGTAAAGC
GTCGTGCTATCTGGACTTATTTAAAGGCTGTCTCAGAAGGTGCCCTGGGAAGATTGGAAAATCTATCTGCTTCTTAT

CTGCGAAATGGCGACCTCTTGTCTCAGTCTTCGTCTCTGTTGCCAGGTTTATCGAGCAAGCTTGG
ATTGCGACAGCAGTGGTAGTTCTCTTAACACTCTTGCTCTCCAGTTGCTGGCCCTCATGCCTTGACTACCAAGTAT
TTGACCCAACCTTTCCGGACAAGGGCGAAAAGAAAAAGCTTACAGGAGGTAGTTGAGGATTGACCAGTTTGT
TACTTGTGGAATTAGTTGGGGTTGATTACCTTCAAGAAAACTAGCCCTCTAGCCTACTAGGAGCTGGTTGGTTTAC
TAGTCTTGATTTGCTTATCAGGTAACGTCAGATGCAGGACTAA

4158.1 (SEQ. ID. NO. 272)

ATGAGAAAATCAATAGTATTAGCGGCAGATAATGCCATCTTATCCTTAGAGACGACTATAAGCTGTATTGTATCACAA
TAGAGATGTGATTITATATTCTAACAGTGATATAAGCTCCTGAATGGTTAAATTATTGGGGAGAAAAATGGAAGTTGTGA
ATTCTACAAATTGCGAGTGTACACATTGATAAAGAACTTTGAAAGCTATAAAACAGGACCTCATATAATTATGCTTCTAC
TTTAGATTTTGCAGACAGAAGTGGTGAATCTGATAGGGTATTGATCTGGATTCCGATATCATTGTAACCTGGGAACTAGC
TACTTGTGAGATAGATCTCAAAGGATATTCAATTGGCTGTGATGATGTCATGCCATAGGACGAAATCTGGAT
TTAATACTGGTATGTTACTAATGGATGTCGAAAGTGGAAACATTCATTGTCATAGTTATTGGATATTAGCGGCCGAG
CAGAATCAAGTTGTCATCTGGGATCAGAGTATTAAATATTATTGAGGATAATTGGCTAGCTAGATAAAACATA
TAATTATATGGGGTATTGATATTACCTTGCTCAAGAATGTAACGCTAGATGACAATCCACCTACAATTGTTCACT
ATGCTAGTCATGATAAACCTTGGAAATACATATAGTATCTAGACTACGTGAATTAGTGGGTTATAGAGATTGGATTGG
TCAGAGATTGCTTCAACGTTCCGATTAAATTATTGAAAGAAGCAATCAGTCTAAAAAAACAAGTGATGCTTGACATG
GAGTGCAGATATAAAACATTAGTATTAGTACACGGTTACGTGATTGCGATTTCATTGGCTGCACCGTGTGATTGTT
CTGAGGAGCTGACCTCTATCACAGTATACGAATGTAACAGTATATCATAAAATGTTACATAGTAAATTGATGGCTATTG
GACGATTCTATAGTTATTAGATATTAAATACAGGTGGAGAGGTTTAATGTAGTTACAAGGGCACAAGAAAGTGGCAAGA
AAATCTCGTTTGTATATCACACGTAAAGTATGGATGATGGACTCTATGACGGTATTITCTGTGGAGAGACCAGATGAT
TTAGTGGATAGAAGAATATAGAGATAGAGTAA

4158.2 (SEQ. ID. NO. 273)

ATGACTAAGATTCTCGTCAATAGCAGTAAAAAAAGGACTATTACCTCATTTCTACTGTATCTATGTATTGGGAAGTCG
TATTATTCTCCCTTGTGACCTAAACTAAAGATTTTAGGAGGTTCAACAGCCTATCTAGCCTCTCACGCCGCCCTAAC
AGGTGGGAATCTAAAGGTTATCAATTCTGTTGGATTATCCCTTGATGTCGCCCATGATTITATGGCAGATGTTTC
TTTTCTAAACGGTTGGTTAACATCTACGCTATAGAAATACAAGATGCCGTAAAATGACCTGACCTGCTAATTGCTG
TGATTCAATCCTGGCAGTTAGCTGAGACTGCCAGTACAATCCTCTATTCTGCAATATGGTTGCTAATGAAATACAATAT
TGCTGATAGCAGGAACATTCTGTTGGTGTAGATTAAATGCAGATGGGGATTGGAGGTCTATTGTAATCCTCC
TATCCAGTATGGTTTAAATATCCTCAGGATGTTGGAACATTCAGACAGTACACATTCCAACAGGGATTATTGTTA
CTTGCTTATTAAACCTTGTCTTCTTACTTGCCCTATGTATCGAGCTCGTATTGGTCTGTAAATAAATGGCT
TACACAATCGATTAAACGCTATTCTTACTCGAACATCTGTAATCCTGAGGTGGGATGCCTTATATGTTATGATGAGT
TTCTTAGTGTACCGCTTATTGTCATCTGGGATTATTCTCTAACATTCAGGGTTAGGGCTTATCAAAGGAA
TTATGGTGGAAAGCCTTGTGGTCTATGTTATATTGGCTTATTAGTATCATTGCTTTGTACGATGA
ATGGAGAAGAGGATTGCAAGCCGTATGAAAAAAATCTGGAGAACATTTATGCTATTATCCAGGTGCCGATACTAGTCGATT
TATTAATCGATTGGCTCTGTTCTCAGTCATAGGGTCTCTTAAATGTTATGGCAGGTGCTCCCATGCTTTGTTTG
TTGATGAAAAGATTATTACGATTGGCAATGATTCTCAGGCTTATTATGATGTTGGGGCATGATTTCAGGATTAGAGACGA
GGTCAAGGCTTAAAGGCTAAATGAGACCTATAGACCTTGTATTAG

4158.3 (SEQ. ID. NO. 274)

ATGTCCTCTTTCGGATCAAGAATTAGTAGCTAAACAGTAGAGCTTCGTCAGCGCTTCCGAGGGAGAAAGTCTAGACC
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GGAGCTATTGTCATGCACTATGGAATGTGCTGAGATGAATACGGGGAAGGTAAAGCCTTGACAGCTACCATGCGCTGTCT
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AAGCTTATCTATGCTCGGATATCATCACACACCAATAGTAATTAGGTTTGTATTCTAAATGATAACCTAGCCTCGAA
TGAAGAAGGTAAGTTTACGACCGTTAACATATGTGATTATTGATGAAATTGATGATATCTTGCTTGTAGTGACAAACTC
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GAGGATTATATCTTAAAGAGGAGAAAGAGGAGGTTGGCTCACTACTAAAGGGGCCAAGTCGCTGAGAATTCTTAGGG
TTGATAATTATACAAGGAAGAGCATGCGCTTTTGTGCTGTCAATTGGTTATGCGATTGAGCTCATAAAGCTTCTACTAAA
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AAGGAGGTCTCATCAGGCTATTGAAGCCAAGGAACATGTCAAATTATCTCTGAGACGCCGGCTATGGCCTCGATCACCTA
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TTGACTATCCAGATAATCTATATACTTACCTTACCTGAAAAGTGTATGTCATCCTGGAGTACATCAAGCAATACCATGCTAAG
GGAAATCTTACTCGTTTGTAGGCTCAGTGTAAATGTCCTCAACTCTATTCGTCCTCTGGTTCTGAAAGGGATTGCCCAT
AATGTCCTAAATGCTAATAATGCCGCGCGTGAAGGCTCAGATTATCTCGACTCAGTCAGATGGGGCTGTGACAGTGGCTA
CCTCTATGGCAGGACGTTACGGATATCAAGCTGGTAAGGAGTCGAGAGCTGGGCTTGTATTGTTATTGGACTGAG
GCGGATGGAAGTCAGCGATCGACCTACAAATTCTGGCCGTTCTGGTGTCAAGGGAGATCTGGTATGAGTAAATTGTT
GTACTCTTAGAGGATGATGTTATCAAGAAATTGGTCTACCTGGGTGATCAAAAGTACAAAGACTATCAGGTCAAGATA
TGACTCAACCGGAAGTATGAAAGGCTGAAATACGGGAAACTAGTCGAAAAGGCTCAGCATGCCAGTGTAGTGTGCG
GTTTCAGCACGCTGTCAGACTCTGGAGATGATGCTGAAAGTATGAAATATACAACGGGATATAGTCATAAAAGAGGAAATCGTCT
AATAGATGGTTCTCGTACTTAGAGGATGTTGTTGATATCATTGAGAGATATACAGAAGAGGTAGCGGCTGATCACTAT
GCTAGTCGTGAAATTATTGTTCACTTATTGTCACCATATTAGTTTGTATGTTAAAGAGGTTCCAGATTATAGATGTAAC
GACAAAAGTCAGCTCGTAGCTTATGAAGCAGGTATTGATAAGAACTTTCTGAAAAGAAGAATTACTTAATCACACATG
ACTTATATGAAACAGTTTACGACTTCACTGCTAAAGCCATTGATGACAACTGGGAGACGAGCTAGACTATCACAACAG
CTATCCATGGCTATGGTGTCAATCTGCTAGTCAGAAAAAAATCCTAAACGCTAGAGTACTATCAAGAAGCCTACGCCGGTTG
AAGCTAGTAAAGAACAGATTCATGCCGATATGGTGCCTAATCTCCTGATGGGGCTGGTTGAGGTACTCCAAAAGGTGAAAT
CGTACTCATTTCCATAA

4158.4 (SEQ. ID. NO. 275)

ATGATAGGGACTTCCGCGCTGCTTGTAGCTGACTAGCAAATTTCATCGTCCCTATTGAAATTACCCCCAATAGTGC
AA TACTGAAATTGCAACCACCAAGATGGGATTGGGCAGGTTCTCAGCAACCTCTGCTCAAACGGTTGACAACCCAGTC
AACGCC CTGCTTACTGCTAATCTTATTTGGGAGTCATTGGATCGTATGAGAGAAGCCAGTAAAATAGTCA

AGAATTGCTAAAACATCGCTGACGTGACTTCTAAATTGTCGAATGGATCATCAATCTGGCTCCATTGGAATCCTGGTC TTGTTTTAAACCATTCGACAAGGGAGTCGGAAGCCTGCCAACTACGGTATTATTGGTCTATTAGTAACGACTATG CTTTTGTGCCCCCTGTGTCACCCCTTGATTGCCCTCTTCTTATGAGACGCAATCCTTACCCCTAGTTGGAACGCTCC GTGTCAGCGGTGACAGCCTTTACTCGTAGTTCTGCGACTAACATCCTGTCACATGAAACTCTGCCATGACCTTGG CTCAACCCAGATACTTCTGTTCTATCCCCTCGGTTACTATCAATATGCCGAGTAGCAGTACCTTAACCTTGG ACCCTTGTGTCAGTTAACACTTGGATTCCCTGTTCTTATCCAGTTGCTTGTAGCCCTTICGGTATTICTAACGATAITGCC TGTGATGCTTCAGGTATTGCCGGAGGTTCCCTCCCTTATCCAGTTGCTTGTAGCCCTTICGGTATTICTAACGATAITGCC ATACAAATTGTTGGGGTTGGTTTGATTGGTGTATCCAAGACTCATGTGAAACAGCCCTTAACCTTCTACAGATGTCCT CTTACCGCCGTTGCCGAATACGCAACCGTAAAAAATAA

4158.5 (SEQ. ID. NO. 276)

ATGTCTATTAGCCAACGTACGACCAAGCTCATCTTAGCTACCTGTCCTGCCCTGCTTGTATTTCTCAATCTTCGTCA GCAGTTTGGCTGGAATTATCGCTCTTGGCTATCTGATACGCTAGAAGTACTTTAAACTGGCTCGCAATCGCTTTTCT TCTATGCTCTAGCTCTGGCTATCGGTGTTCTAGCTTTACTGAGCGGATTTCATATCTGGAGTCTGCCCTATCTGGCC TTCTACGTTCTTCTAGCTACAAGATGGGCTGGAAATTGGCCTACACCAAGCAGTGTGTTAGCCATCTCTTGGTCA AGAGTCACCTCTCAGAGCTTCTAGTCATGAATTCTCTTCTTGTATTGGTACAGGATTGCTTGTGTTAATCTTA TATGCCCTCACGAGAAGAGGAATTACGACTACACACGGCTGGAGAGAAAAGTTAAAGATATCCTCAGCGCTTCAA ATACTATTATCCAGAGGAGACGGACGCAACCGAGCACAGCTGGTAGCAGAATTAGACACGCTTGTGAAAGAAGCCCTAG ACTGGTCTATTGGATCACTCTGACCACCTCTTACCCAGACAGACTACCATATCCACTACTTGTAGATGAGACAGCGACAAA GTCGTATCTGAGAACATGGCCAAACAGATTAACACTTGTACCTTGCCAGTGAAGGCTGTATCTAGCGCAACTTTTCTC AAAAATTGCAAGTCAGTACGAGCCAGAACATCCTGCTTGTAGATGAAACGTTATCTGGAGTCTCAGCGTAAAGCCAAACCTTC GAACCGCAGTCTGCCCAAGACAAGAGAAGAATTGAAACCCGCCACCCCTCTCAACTCCTACGTGAAGCCAAACCTTC ATCCAAGTAAAAGTTGATTACCAAAATATAGACAGTAA

4158.6 (SEQ. ID. NO. 277)

ATGGAAATCATGTCGCTGCGATTGCTGTTTGCCTCATCATTGGTTAGTCATTGGATATGTCAGCATCTCAGTAAGATG AAATCATCTCAGGAAGCTGCAAGAGTTGATGCTTTAAATGTCGAACAAAGCAACTAATTACGTGGACAAGCTGAGCGT AAGCGGATTACTTGTAAATGAAAGCCAACGTGAAAGCAAGTCTCTTAAAGAAGCACTATTGGAGGCCAAAGAAGAAG CCAGAAAATACCGTGAAGAAGTGGACGCTGAATTCAAAATCAGAACGTCAAGAACACTCAAACAAATCGAAAGCTGTTGACAG AGAGAGCTACTAGCCTTGACCGTAAGGACGACAATTGACGAGTAAAGAACAAACACTGAACAAAAGAACAAAGTATT CTGATAGAGCGAAAACCTTGTGCGCTGAAGAGCAATTAGAGGAAGTCGAAGAGACAAAAGAACAGAAGACTAGAGCGT ATTGGTGCCTGTCAGGCAGAACGAGGATATTATCTGGCTCAGACAGAGGAAAATTGACAGGGAGATTGCCAGTC GCATTCGCGAAGCTGAGCAAGAGGTCAAGGAAGCTTGTGACAAAATGGCAAGGACATTGGTCAAGCTATGCAACGTAT CGCTGTAATATGAGCGAGTCACAAACTCAACAGTTCTGACAGACGATACTATGAAGGGACGCTTATTGGTGT GAAGGTGTAACATTGTAACCTTGTAAAGGTTGACAGGGGTCGATGTGATTATGACGATAACCCAGAAGTGGTGTACCTGT CAGGATTGATCGATTGTCGTAAGGTTGCTGACTATGAAATGTTGCTAAAGATGGTGTATACATCCAGCTGTG ATCGAAGAGTGGTGAAGAAAACCGTCAAGAGATGACAAATAGGCTGAAGGCTACATGGTGGAGGCTGTGCTTATGAAATG GTGCGCCAAAACCTTATCCAGACTTGTGAAGGATTGGACGTTGCTGAGTCTGACTTATGGACAAAATGTTGCG CATTGAGGTTGCTAAGTGGCTGTTATGGCAGCGAACCTGGTGAAGGTAAGGCTGAAATGGTGTGCTTGGCGTGTGCTGATT CCTCACGATATGGAAAGCCATTGACCATGAGGTTGAAGGTAAGGCTGAAATGGTGTGAAATGGCCTGTAAGTAC AAGGAACCCCAGTTGTTGTAATGAGATTGCTAGTCACACCGAGATGTTGAAGGCTGAGAGCTGATAAGCAGTTATGTCG CTGAGCAGATGCCCTGAGCGCAGCCGTCAGGTGCTGAGTGAAGTCTCTGAAAGCTACATCAAGCTCTCCATGATTG GAAGAATTGCTAACGGCTTGAAGGAGTGCAAACCTAGCTTGCCTTCAAGGAGACGCTGAAATTGCTATCATGGTCAATC CAGGAAAATCAAGGAGCACAAAGTCACAACTTGGCTCACAAAGTGTAAAGAAAATTGAAACAAATCTCGATTATCCAG GAAATATCAAGGTAACCGTGAATCGCAGCTCGTAGATTGCTAAATA

4158.7 (SEQ. ID. NO. 278)

ATGATGTTAAAACCTTATTGATACCTTGTGCGACAAGGTTCTCAAAATATTACCTGTAATCTGGAAAGCAAAACGTGC CCAGCAATTGGAAGCAGGTGCCAGCAACTCAAGGTTCAAGTCTGAAAAATCAACTCTCGCCTTGAAGAAGAAATCGAA TCAGGAAACGTTACAATTCAACCCAGATCCAGAACAGGAAAACGTGAAGCAGTGCCTGCCGTATCGAAGAAGAAAACGCCGC AAAGAAGAAGAAGAAAAGAAAATCAAAGAGCAAATTGCTAAAGAAAAGAGATGGTAAAAAATTAA

4161.1 (SEQ. ID. NO. 279)

ATGTCAGCATATCAATTACCGACCGTATGGCAGGATGAAGCTAGTAATCAAGGGACTTTACGGGCTAAACAGACCAACAG CAGGTGCCCTTCAACAAAACCTGCCAAAAGGAGAACAGCTTTCAGCTTATTACTGGAACACCAAATGGTGTGAA GGTTACTATCTTATTGGAAGAATTACTAGAACGCTGGTTTAAGGAAGCGGCTTACGACTTGTATAAGATTGCTATCATGGATG GGGATCAATTGGATCACAGTTGTGAAGCTCAATCCAAGATTCCAGCCTTATGGACAGTCAGGTACTGAAA CGTAAGACTTGTGACTCTGCTCATATTCTTCTACCTTGTGAGAAATTGGACGCTTACCAAGTAATCCTGTGAAAG GGTAGAAGGTTGAAATTGGTATTCTGGCAAGCAGGTGACGACCTTCTAGGTGGGGATTGGACATTCTCAATTATG CTCTGAAAATTGGAATATCTTATTACCGTTTACGATGGAGTGAAGCTGAAACGCCAGTGGATTATTGGATAAGGAATTGGCT CAGAAACCTTATATTGCAAGGAACTAGACTACAGATTGACGATATTGCTATCTGGTCTGGTATGGACAGTTAGTTCAAGGAA ATCTTACCAAGGTTCTGCAAATTCTGGATGCCCAAGTTATCAAACATCTAGAAAATGGCAGAAAAATTGCCAATCG TCCAGCTGTTAGCGTGGCTTGGAGTAACCTTACAGAAATTAG

4161.2 (SEQ. ID. NO. 280)

TGGCAAGCTTGTACCTTCTATCATCATGTTCTATGTCGGTTTCGATGTTCTAAGAGATACCAATTCAAAAGATTCTCAGTCGG GAAGAAACGGTATTGATCCCTTGGTCAACTCTAGGAATCATCTCAGCGATTATGTTGCTATCTCTACAATAC TCGCTCTAGTAAGAAATCCAACCTCAATCGCCTGAAGGCAGCTGCTAAGGACAATCTTCTGACGCTGTTACCTCACTTGA ACCGCCATTGCCATCTAGTAGTTCAATTACCGATTGTGAAACTGGTTGCTATCATCATCACCTTCTTATCTG AAGACTGCCTATGATATCTCATCGAGTCTCTTCTAGTGTGCTTACCATGAGTGGCTTACGGTACGCAACATCTACCTGGTATT GGCTATCATGGAAAATTCCAAAATCAGCAAGGTCAAATCGCAAGGAGGTCGACCTACGGTAGCAACATCTACCTGGTATT AACTAGAGATGAATCTGACTTGTCTGTTGAAAGCCATGAAATCGCGATCAGTCAGTCTATGCTGGAGGAGCGTT TTGGCGTCTTGTATACCGATGTCCATATCGAACCCAGCACCTATCCCTGAGGATGAAATTAGACAATGCTATAAAAATTG CTTATGCGTGAACAATTGATTGACCAAGGAAACCAACTAGAAGAACTCTGACTGATGATTGCTATATTGCCAAGATG GAGAGCAGATGGATAAGAGGCTTATAAGACCAAAAAAGAGTTAAATTCTGCTATCAAGGACATTCAAATTACTCCATCAG

TCAAAAAACCAAACATCTGCTATGAGTTAGATGGTATCCATACCAGTATCTGGCGTCCCACGAAACCTGGAAAAT
ATCTTCATCAAGAAACAAAAAAGAATAG

4162.1 (SEQ. ID. NO. 281)

ATGACAATTAAACTAGTAGCAACGGATATGGACGGAACCTTCTAGATGGGAATGGACGCTTGTATGGATCGTCTCAAGT
CTCTTGGTTTCTACAAGGAAAAGGGATTACTTTCGGTAGCTTCGGGTCGGGATTCTGTCTAGAAAAATTATT
GCTGGTGGTCTGATGACATTATTCATCGCGAAAATGGCAGTTGGTAGAGTATCAAGGTCAAGGACTTGTATGAAGCGA
CTATGTCTCGTACTTTATCTGGCAACTTGGAAAAGCTGAAAACCTCACCTTATGTAGATATCAATAAACACTGCTTGCAG
GGTAAGAAGGGTTCATATGTTCTAGATACGGTTGATGAGACCTATITGAAAGTGAATCAGCACTATAATGAAAATGCCAA
AGTAGCGAGTTGGAAAGATACAGATGACATTTCACACCAACTTCACAGAAGAAGCCTGGAAAGATGGG
AGGCTGGGAAACGAAACCTGGTTAACGGCATGACAACCTGGCTTGAATGGCATTGATATTGTTCTGGACTATGTC
GATAAGGGAGTGGCATTGTTGAATTAGTAAACACTGGTATCACAAATGGATCAGGTATGGCTTTGGAGACAACTTAA
ATGACTTACATATGATGCAGGTTGGGACATCTGTAGCTCTGAAATGACGACCTGAAATTAGAATTAGCAAGAC
TGTGATTGGTACCCATAAGGAACGGTCGGTTAGCTTATGGAGGGCTTATAA

4162.2 (SEQ. ID. NO. 282)

ATGGCAGATAAAATTGATTGCTTGGACTTGACGGGACCTGCTGACTACTGATAAAAGGCTGACGGATCGTACCAAGG
AAACCTTGCAGCTGCGGTGATCGTGTATCAAGGTCTGTTGACAACTGGTGTCCCTTAAAGGCCATGGATTCTTCTC
CATGAGTTAGGGACTGACGGTCAGGAAGATGAGTATACCATTACTTTAATGGTGGATTAGTTCAAGAAAATACAGGAGAAA
TCCTTGATAAAACAGTCTTTCATATGATGATGTTGACGGCACGTTGTATGAAGAAACAGAGAAATTACTGCTCTTGATGCC
ATCTCAGAAGGAACAGTTCAATCCAATCGGACCAAGAAAGTCTTATGCCAAATTCACCCAGCTTGGACTTGTCC
AGTGGACTTTGAAGACTTACTGCTAACACCTAACAAATGCGTACTGCCCTTGTCAAGAACCCCTGGATGCCAGCC
ATTCAGAAAGATTCTCCAGAATTGTTGACCAATATGAAATTTAACACGTTGAGAATGGTCTAGAATGGTACCAAAGA
ATGTTCATAAAGCAACAGGTTGGCAAAACTAATCAGCCATCTGGATGACCAAAGTCAAGTGAATGGCTTGTGGTACGA
GGCAATGACCTCTCATGATTGAATGGGACGGTCTGGTGTATGAAACACGCTTGTGAAAGTAAAGGCAGCGCA
AATGTTAGTGACGCCGATGCCAACGATGAGGAAGCTGTCGCCCTGGCTATGAAAGAATATGTCTAAAGGAGAACTAA

4164.2 (SEQ. ID. NO. 283)

ATGGAAGATTACTTATTCTTATTAAATTGCCAATCTAGCTGGTCTCTTCTGATTGGCAAGGGCAGGATAGGCAGGAGAA
ACACTTAAGTAAGAGCTGGAGGATCAGGCAGATCATTTGTCAGACCAGTTGGATTACCGTTGACCAAGCCAGACAAGCC
AGCCAGTTAGACCAAAAAGATTGGAAGTGGTGTGTCAGCGACCGTTGCAAGAAGTGGGATTGACCAAGGCTGACGAAC
CCAAGTCCGTCAGAAATGACAGATAATCTCTCCAAATAGAGACAGACAGACCAACGCTTCAAGGCTTGGACAGGAAC
AAATGAGCAACGTTGGAAACAAATGCGCAGACGGTCAGGAGAAAACATAGAAAAGACCTTGCAAGACAGCCTTACAGGCTT
CTTGGAGACAGATTCTAAACAACCTGGAGTCTGTCATCTGGCCTTGGAGAAAATGCAAGACAGTGGCTGATGTCGAGGCT
CTTAACAAGGTTCTCTGGAAACCAAGACCGAGGGATTCTGGAGAATTGCAACTGGGCAAATTATTGAAGACATCATGA
CACCTGCCAGTACGAACGAGAATACGCAACGGTTAAAACCTAGTGAACGAGTGGAGTATGCCATCAAGTTACCCGGAC
AAGGCAGACAAAGAATACGTCATCTGCAATTGACTCTAAGTTCCACTGGCAGGATTATTACCGCTTGGAGAAGCCTATGA
GACAGGTGACAAGGATGAGATTGAAACGCTGTCGTAAGTCACCTGCAAGCGCTTGGAGGATTAGGGATATTAGGAAC
AAAGTACATAGCACCCCTCGGACGACCAATTGGAGATTGTTGACAGAAGGTTCTACTCAGAAATCGCCCAA
TCGGCTTCTTGTATGATTGAGACGGGAAGAACAGATTATGTTGACGGACCAAGTACCCATTGACGCCCTTAACTCCC
TATCAGTTGGTTCAAGACCTTAATATCCAAAAGAGTGGCAGCATATCAGCAAGACTCTGCCAGTGTCAAGACCGAGTT
GGCAAGTTGGTGTATTCTGGTCAAGGCACAAAACATCTCAACATGCCCTGGCAATTGATGAATTATTAAACCGTC
GTACCATAGCTATGAGCGGACGCTCCGTACATTGAGTTGTCAGAAGGTGAGCCTGCGCTGATCTACTCCATTTCAGAA
AATGAGGAAGAATATGAAGATTAG

4164.3 (SEQ. ID. NO. 284)

ATGAAGATTAGTCACATGAAAAAAGATGAGTTATTGAAGGGTTTACCTAATCAAATCAGCTGACCTGAGGCAAACCTGAG
CTGGGAAAAAAACTACCTAGCCTTACCTCCAAAGATGATAGTGGCAGGATTGATGCGGAAGCTCTGGATGCCAACCTCATAA
CATTGAGGCCCTTACCGCAGGTAAAGTTGTCACATGAAAGGACGCCGAGAAGTTATAACAAATCCCTCAAGTCATCAA
ATTACTCTCCGCTGCCCTCAAGCTGGTGAACCCAATGACCCAGCTGATTCTCAAGGTCAAGTCACCCAGTTGATGTCAGGAAA
TTCTGACTACATGTCGAAATGATTTCACAAATTGAAAATCCCTGTCGGCAACGGATTGTCGGAAATCTCTACACCAAGTAT
GATAAGGAATTCTACTCCATCCAGCTGCAAGACCAACCCACCATGCCCTTGAACGGGCTTGGCTATCATACGGCAGCA
TGGTGCCTTGGCAGACGCTATTAGCGAAGTTATCCTCAGCTCAATAAGAGCCTGCTCTATGCCGGGATTATGTTGATGAC
TTAGCTAAGGTACGAGTTGACGGGGCCAGACGACAGAGTACACAGTGGCTGAGGAAATCTCTGGACATATCGCTCTCA
TTGATAGCGAAATTACCAAGACAGTTATGGAACCTGGCAGTCAACAGGAAAGTGGCTTGTCTGTCATGTCAT
CCTCAGTCACCCAGGCTTGTGAGTATGGAAGCCCAGTCCGTCACCGATTATGGAAGCAGAGATTATCCATATGATTGAC
AATCTGGATGCAAGCATGATGATGTCACAGCTTCTGGGATAAAGGAGAGATGACCAATAAAATCTCGCTA
TGGATAATCGTTCTTCTATAAACAGATTAGATTAA

4166.2 (SEQ. ID. NO. 285)

ATGAGTAAAAAGCTAAAAAGGGTTAAGATGCCCTCATCTTACACCGTATTATTGATAATCATTGCTATTATGGCAGTGCT
AACTGGTTTATCCCTGCGGGGCTTATAGAAGGTATTACGAGACTCAGCCTAAATCCACAAGGGATTGGATGTC
TCATGGCACCGATTGGCTATGCTAGGTACTCATCCAGAGGAAGGTTGCTCATAAAGAAACGAGGCCAGCAGATTGATGT
AGCCCTCTCATCTTATGGTTGGTTCTTGGCATTGCAACAAAACAGTGGCTCTGGACGTAGGGATTGCCCTATCGT
GAAGAAGTATAAGGGCCGAAAAAAATGTTAATTGGTACTGATGCTTGGGTTACCTGACTGCTCTTGGTACAACCTTATGGTATGG
GTGAAGAACAAATGGCCCTCATCCACTCTGGCAGTTGACTGATGGGGTTGGGTTACCTGACTGCTCTTGGTACAACCTTATGGTATGG
ATTGGTCTGGTCTCAATCGGCTTGGCATCTACTCTGAATCCATTGCGACAGGTTGCTCAGGACTGCGGGAGTT
GGTACAGGGGACGGTATCGTACTCTGCTGATCTCTGGGTTACCTGACTGCTCTTGGTACAACCTTATGGTATGG
GATAAGGATTCAAAAGATCCGACTAAGTCAGTGGTTATAGTACTCGCAAGAACAGATTGAAACACTTAAACGTAGAAGAAT
CTTCATCTGAGAATCTACACTAGCAGCAACAAAATGTTAATTGGTACTGCTCTTGGTACAACCTTATGGTATGG
GCTTCACTCCATGGGACGACCCCTGGGTTACCTTGGTACAACCTTATGGTACTGCTCTTGGTACAACCTTATGGTATGG
GTGTTATTATGGTCTTAAAGAAGATAAGATTATCTCTTCTCATGAATGGTGTGCTGACTGCTAGTGGCTTGTGATCG
TAGCGATTGGCTGTTGATTCAAGTTATGAAACGACGGTATGATTACCGATAACCTCTCAACTGGGTAAGAAGGCTT

GAGCGGTCTATCTTACAAGTCTTATCGTTGAACTTATCTTCTATCTACCTATGTCATTCTGATCCCATCTCATCTGGT
CTTGCAGCGCAACTATGGTATCATGGC
TCACATTGGAGAATTGAAATGTCGCTCTAGCTGATTACTGCTTACCAACTCTGCTTCAGGGTCTTGAACCTGATTG
ACCAACATCTGGTATTGTGATGGGAGCTTGACCTGGACGTATCAACATTGGTACTTGGTGGAAATTATGGCAAACCTG
TAGTCGCTATTATGTAGTACCATGCCCTCTTCTCCTGGAACCTCCATTCTATAA

4166.3 (SEQ. ID. NO. 286)

ATGAAAATAGATATAACAAATCAAGTTAAAGATGAATTCTTATATCATTAAAAACCTGATTCTTATCTTCAGTACTCAA
TGAAGGAGAAAATGGAACACCTTGGACAACCAATCCAAGATGTCCTAGAAAAAACTTAGAGATTGTCGAGACATAGG
TTTCACTACCTATCTTGCACCTAAAGGTATTACGGATATGCAAGAAATCGTCAGGGAGCAGAGCTCTGCCATTCTGTC
ATTGGATGTTGTCATCAGGTGATGAAAGCAGATTGGCAGACACCCGCTTGAAGCAACTATCAAAGACGGCTGGTATT
CGGACGGTGGTCAAGATGAAAGGCGCTTGCCTGAGCTCTATGCAAGTAAAAGCTGTCAGGACCAAGGTATTGAG
TTCAAAAAGCGCGTACGCTTATCTTGGTACCGATGAGGAAACCTCTGGCGTGCATGGCACGCTACATAACCATCGAAG
AACAGGGCAGTATGGGCTTGCACCTGACTCATCTTCTCTGACCTATGCTGAAAGGGCTTACAGGTCAAACCTCAT
GCCCTGGATCGATCAACTAGAGCTGAAAGTAGGAGGCCCTTAACGTTGACCTAGACAAGGCCACTACCAAGGTCTCC
TCTATGAACAGGTTGTAACGGTCTCAAAGAAGCTGGTTATGATTACCAACCAGTAACAAACCGTAACGGTCTCGGAGT
GCCAAAGCATGCTAAGGATGCTAGCAAGGTATCAATGTCATCCGACTAGTACCCATTGCTCTCCAAGAACACC
CTGCTCTCAGTTCTGCAACACAAGCGTCAAGCGGCACAGGAAGACAATCTTGGTGTATAGCAGATGAAACCTT
TGGTACCTATCTTAAATGTCGAGGTCTCATGATCAATCATGCAAGCTCTGGTATTGACATTGGACTCTGTCTT
AGCTGACAAGGAAGAACTAGTAGAGTTGCTTACAAGATGTGCAAAAACCTACCAACTCCGCTACGAAGAGTTGACTATCTA
GCCCTCTATACGTCGAGAACAGTAAACTGTTAGCACACTGATGCAAAATCTACCAAGAAAAGACTGGCGATAACAGTC
CTGCTATTTCATCCGGTGGTGCCTTTGCTCGCACCATGCCAAATTGTGAGCCTTGGCCTTATCCAGGAGCGAAG
CAGACAGAACATCAGGAAATGAATGTGCCGTTAGAAGATTGACCGTCTAGGATATTGCGAACCGTCTATC
GACTTGCAACTTAA

4169.1 (SEQ. ID. NO. 287)

ATGTCTAATTCTATTGTCAGTTAGTCTCTCAATTATTGCAAATTAGCAGATATTCTTAGAGTAACAATCATTGCT
AACATATACATTATTCAAAATCAGTAATTGCCACATCACTAGTCTCTATCTTAATAGGAATATCCTTTGCGAGCTT
TTAGTCTCGTGTACTAAAGGTTAGCGCTAACTTAAAGGGTTTATCTTCTCTCAATTGGAAAGACTATATTGGCGAT
ACTGGTAGGAATGTTACCGTAATGCAATCCGTAGCGCCTTGGTACCTATCTTGTGTCATGGCAATTCCATACTAGATG
GTTTGAGCACCCGTTCTATGCTATTGTCACGCTATGCAACCGATTGGTAAGGCTAATTGACCTTATCAATGACT
GGTGAAGCTTCAATTGATAGGTTGGGATTAGGTTGACTCTTGTGCAACAATTGGTCTTACCTACCGTGTATCAA
TTAGTCTTGATATCATTTCTAGTTCTGATGTTATTCTCTCTAACGCTGAAAGTGGAGGTGTTAGAGTCAGAAACTAATCT
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AACCCCAATTATTCACCCCAATCTAAAACCATCCAGAACCTTGCCTAGCTTAGATCTGGATGGTTCTTACCC
ATGGGTGTTTACTAGACAAAAAAGAGTTCCCTTATGGTATAAGTGTAGAAAAAAACACAAAAGAAAGGAAACTCA
CATGAAACAGTTACCAATCATCCTTACCCAAAACAAGTCTTTTACCAACTATCTTGCATGGAGGTCAATTAAACCCAGTATG
GTGGTCTTATCTTTCAGGAACTTTTCCAGTTGAAACTAAAAGAGCGGATTCTAAGTATTAGTAACGAATGCCAA
CGCGCTACTGCGTATTGCGATTGAGATCTTGTCCAGTTCCCTTCAACTGTTAACAGGTTATGAAACGGACTATGCT
TGAAAGAATTGTCAGCTGATGCCACTT
TCCAAAATTGTTGAAAGGGCAGCTTGCCTCACAGCCAACCTTATCCGTTCTTCCAGAACTGACGGAGGAAACAGTCC
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GCTTCGAGGGAAAGACAGGTTATTGTTCAATGCCAGCTCTGTTGTAATCGTATTGTTCTGAGGAGCAGCT
TATCACACCTGTTAGAACGGTTAATCAACTCTTCTGCAATGGATAGTGGCTTGCACCCAAAATTATACGATTAA
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CCACAGGATGAGGACTTACCATCTGCCACTCCGCTACTCAGAAACTCTATCAAGCAGGATCTGGTCGACAA
GTCGTGTCGCCAGTTCTGAAACGAAAAGAGGAAACTGTTCTACGATGTTCTCAGTACAAATATGACGAGTGG
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GGGATAAAACGGATAGTCAACCTTAACTAAAACGAAAGTCTGATGAGCTGATCGCTACAACTCTATCTT
CTCAACATCTAGCTGGAGGTACTTCAAACCTTAAACATCAAACGCTCCGCATCTTCTCAGCTGGTGGAAAATG
TGGTCAACAGGACGCAAGCAGCTCTCAAATTGCTAGTCTATGCTTACCTGCTTACGGACTTATCTAGGAT
TAGAAAAGTCACCTGAATCTCCTGTTCTATGAAACCACCTAGAAGAAAAGCTGTTAATGATGCTTAA

4169.3 (SEQ. ID. NO. 288)

ATGATGGAGTTTCAACAGCTCCTCATTTAGAGCCATATGGCAATCCTCAGTATTGTTATGTCATTGCTGCAACCTTG
CCCATCTTATAGGTCTTTTCAAGAAACGCTTGCCTGGTATGAAGTGGTAACTCTCTTCTTATGTCACCATGTT
GTGGTGGAAAGACCAATCAACTAGTCTGCTTGGTATTACCTTCTGCTGGAAATTGTCCTCTTCTACAAGCATT
TCGAAAAGCAAGGATGCCAGTCTGCTTGGTCTCTACTAGTTAGTCTGCTCTCTCTACCTCCGATTATCTTGTCAAGGTGCAAC
CAGCTATCAATGCCAGCTTGTGGTCTTGGGAAATTCTTACCTGACCTTCTGCTGGTGGAAATTGTCATCGAC
TGAGAGATGAGTGTAGGATTACGGATTACCCCTCTGGAAATTCTCCGTTCTCTCATGCCAACTTCTGAGTGGCTTCA
TCGATGCTTAAAGCATTAAATGAAAATTATCAGGCTATCTGAGCGAGATGAGTTGATGGAATTGCTGGATGAATCTGTC
CGCTATATCATGTTGGGTTTGTATAAGTTATCCTAGCTCATGTTAGGAGAGACCTTACTACCTCTCTGAAGAATT
GCCTGAGCTAGGTGGCTTAACTCTATGCCCTGGCAGTTATGTTACTTGTGACTCTTGTGACTCTTGTGCA
GGTTACTCTATGTTGCTTGGCAGTCTCAAACCTGATGGGAATTCTGAGCCCTATCAACTTAAACAGCCCTTTTATCAAGG
GATTTAAAGGAGTTGAAATCGCTGGCATATGAGTCTGCTTCTGGTCCGTGACTTGTCTTATGCAATGGTATGGT
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GGCATGGTGTGACCTGGTACTATATGCCCTATGGACTCTTCTCATGGACTAGGCTGGTACATGATGCCCTGGTTGCAAG
AAAAAAACGCTCAATAAGGAACGGAAAAAGCAGGGAGGGTACCTGAGAATCGCTGGATTGAGCTTGGCAT
GTTGTCACTTCCATGTTGTCATGTTGATCTTAATCTTCTGGATTCTGAAATAATCTATGCTTAAATA

4169.4 (SEQ. ID. NO. 289)

ATGCTTAAACGTTATGGATGATCTCGGACGGCTTGTGCTGGTTGGCTTCTGTCATTTCTTATCCTACTG
AGATGCATATAATCTAGGAGCTAAAAGCGTCAGCAGTGGCTACTACTATCGATAGTTAAGGAGCGAAGTCAAAAGT
CAGAGCACTATCTGATCCAATGTGCCCTTGTCCCTCTGCTTAGTGAATGGCTTGCAGGGTGCATCCTGC
GGTATTAGCTGAGAAATACAATCGTCCTACCGTCCTTATCTTCTAGGACAGGGGGAGCTGCATCGCTAACCAATATTTG
GAATGCAACAGATGTTACCACAGCTGGAGAATAAACAGTGTATGTTATCTCACCTCAGTGGTTCACTGGGATCAGGCTAGTC
TGATCCAGCAGCCTCCAGCAGTATTTAATGGAGCACCTGACTAGTTCTGAAACATCAATCTGGGATCAGGCTAGTC
AAATGCGAGCAGCAGTACAGTCAAGTGGGAGCTGAGTAAAGGAGCTGGTCAAGAAGTGGCAAGTAAAGAAGA
ATTGTCGACAGCAGACAATGAAGATATTGAATTATTGGCTCTTAAATGAACGCCAGCTTCTTGTAGTTCGG
TAGAGGCTATGTTAATCGATAAGCATGTAAGTATTAAATCTGCCAGACCAGTCTTCTTATCAGGCAATAGAA
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AGAAGGATTGAGAAATTAAAGGATTCTCAGAAAAGCTTACCTATCTCACTGCCAGAGTATAATGACTTGCAGTGGT
TTAACACAGTTCTAATCTAAGGAAACCCGATTTCATCTCACCTGTTAATTTAAATGGATGAACTATGCTGGT
TACGAGAGGATGATGACCAACAAACGGTGCAGAAGATGGCTTACAGTAGGAAAGTCAGGTTTACCAATATGAGCAGATT
TTCTAAGGACGGCGGGAGCTTCTTATGAGGACACCATTACCTGGTTGGGGTTGGTTGGCTTGCACAGGAG
TGATCTTCTTATCCAATCCCACACCAGCTCGACTTACCATCTGAATGAGCGCTTICAGCAAAGATGGCGACTTAT
GATGGAGATGTCAGAAATTCAATAG

4169.6 (SEQ. ID. NO. 290)

ATGGAGAAAACCTCAAGGCTTGAACACAAACACAGACCAAGAAGGCCAGCAATTGAACCTGAAAAGGCAGAGGATAC
CAAGACAGTCCAAATGTTACTTCGAGGATGCAGCTGCAAGGACCCACCTTGAGTACTATGCAGGTAACTGGCAATCA
GTTTATCCTTCTGAAAGACGGCACGTTGACCAAGTCTTGACTACAAGGCTAAGTGTAGCTGTAAGATGACCCAGGCTGA
GTACAAGGCTTACTATACAAAAGGCTATCATACAGATGTGACTAAGGATTAACATTACTGATAATACTATGAAATTGTC
GGTGGACAAAGCAGAAGAAATCACTTACAAGTGTGTTAAGGAAATTGACTTACAAGAAGGCAATCGTGGCTGCGT
TTCCTCTTGAAGCCACAGATGCTGAGCTGGACAATTCAAGTATGTTAGTGTGACCAACATGTTGCCAGTTAAGG
AGAACATTCTCATCTTGGAGGACAAGCCAAGGCCCTTCTGAAAGAATGGACAACCTGCCACCTACTACCA
GATAACCTATCTGCCAAGAAATCCCCAAGAAATGTTGCCATTGA

4170.3 (SEQ. ID. NO. 291)

ATGAAAGATGGTCATTGCTAGCCCACATATTGTTGAAATGGGGGATTTCAAAAGTTACTGAGTCAGATCCTGA
GGCTTITATAGGGGTGAAACAGGGCAAGATTAGCGTTTATGAAATAGTGAACACTGGCTGCAACTGCACAGATATC
GGCCTGCGACTGGACTTGCATAATACGCTGACGACTATGATAAAAAGCTAGAGGAACAAAAGCTTGAATTGTTAGTC
CGTGTGGAAAAGACAAGCGTAAGAAGTATTAGTTAACGGAGTTAGGCAAGTCCCAGAAAAGAAGTGGGATCGTGTCA
GTCAGAAATTGGACTATCTTACAAGGATTTCAGAGGAAGAAATTCAACATTGAAAGGTTTCAAGAAAGAATT
GGCAATCTGAAAGAGAAGGAAATGAGGTTAG

4170.4 (SEQ. ID. NO. 292)

ATGACTAATTAAATTGCAACTTTCAGGATGTTAGTGAATTGTTGACAGCTCTATCTAACATTTGAGTTGCTGCTTITG
ACCTTGTACTAGCTATTGCTTGCATTCCCTGGCTTTCTCGCTATCATGAGAAGCTGGCCGACTGGGCTTGCAG
ATTGCAAGCTTCTAGTGTATTGCTTGCATTCCCTATTGCAAAATACTACTGGCTGAAGGAAATTGATCCGAACCTGCA
AGAGCAGCTGGATTGCTTGGGATGACCAAGATGGGACGCTCAAGAAATTGAAATTCCACTGCCATGCCATTATCATG
TCTGGATTGGACGGCAGCTGTTGATTATCGGTACGGAACCTGGCGCTTGTGATTGGTCAGGGGACTAGGTTCTT
TATTCTTGGGATTGACCGTAATAATGCCAGTTGATTGGGACTTCTCAGGTTTGCCTTGGCTTAAATTACTGGGCTGT
CTTCTACTAAAAGTGATGGAAAAGCAAAATTACGGACGATTCTCAGGTTTGCCTTGGCTTAAATTACTGGGCTGT
CTTATAGTCCAGCTTGGGATGAGGAAAAGAGAAGGAAAATTGCTGATTGCAAGGAAATTAGTCCAGAAACCAGAAATT
GGCAATATGATAAGTTGCTGATTGAGAAAATACCAGCATGACTGCGACTGTTAACCGAATTGGGAGACAAGCTT
CTTATGAAGCTCTGAAAAGGCCATATTGACATCTATCTGAATTACTGGTACGGTACTGAAAGTTGCTCAACCATC
ACCAAGGTAGTCATGAAACAGAGTTTACGGTGGCGCTGATGGCATTGCTAAGCAGGATCATCTAGCCTATCTC
AAACCCATGTCTTACAAACACCTATGCTGAGCTGTTCCGAAAAGATTGCTCAAGAATATGGCTGAAGGACATTTCAG
ACTTGAAAAGTGGAAGGGCAGTGGAGGCAAGGCTTACCTGAGCTTAAACGACCGCTGAAGATGGAATAAGGGCTTGC
AAATCAATGTATGGTCAATCTCAATGTAGCGACCATGGAGCCCTTCGCTATCAGGCTTACAGTCAGGGGATATTCAA
ATCACGGATGCCATTGCACTGATGCCATTGGAGCGTTATGTTACAGGTCTGGAAAGATGACAAGCAACTTCCAC
CTTATCAAGGGGCTTCACTCATGAAAGAGCTTCTCAAGAAACACCCAGAGTTGGAAAGAGTCTTAAATACATTGCTGG
TAAGATTACAGAAAGCCAGATGAGCCAGCTCAACTACCAAGTCGGTGTGAAAGGCAAGTCAGCAAAGCAAGTAGCCAAGGA
GTTTCTCCAAGAACAGGTTGTTGAGAAATGA

4170.5 (SEQ. ID. NO. 293)

ATGATGCATACTATTGCAAAAGAAAATTGAAAATATCAAACACCTAGGTGAAATGTCAGGTGGTACCGTCGTATGG
TTGCGGCTATGGCTGATTAGGATTTCAAGGAACTATGAAGGCTATCTGGGATGACCTCTTGGCCATCGTAGTTTGGCCAG
TGATTTATTGCTGGTTAGGAAGTTCCTCTGGCTGGAGTTGTTACGAACATCGTATTGTCAGTGGATTGGGATG
ATTGTAGCTGACAGGGATTATCTGTAATCTTGTATCGGAAGGCTGAGCAAGTAATTACTCTTGGCTTGAATTACT
GTATTACCTTATTGGCCCTACAGAAAGCTTATGGTGAAGGCTGAGCAGTACTTACAGTCATGCAGGCAAT
TGACTCTAGTTGGATTATCAGGCACAGTTAAGAAGGAAAAGCAGGAGTTGCTGCCGCTAAACTGACCGCAAGGGC
TGGACAAAGTATCTTCCATTAGTGTGTTGGGGTTGGCTTGGCTTATCAGTCTATTGGTGCCTACAGTCAGGGGATATTCAA
CGTGTATTCAACACAGATGCAACCAATGGGGTAGGGCAAAATCCTCATGACAGCTTACCGTGAACAGTGGATATTCTGG
CGGCTACCAATGCTTCAATCTCTGGGGAGAAAGCCTGCAAATCAAGGAAATATCTAATTATCTCATTAAAC
AGTCTAGTTGGTGGTATCAATGGAGCAAGGCAAGCTAAGCAGAATACTGATTACTTAACAG

4170.6 (SEQ. ID. NO. 294)

ATGAGAAATATGAGGCAAAATATGCTGTTGGTGGCTTTCTTAAATTGACTTATGCCATTGTTGAGTTATTGCAAG
TGAGTATTGGCTAGCGCTGTTGCTGACTCTGTGATGACTTGGGAGATGCGATTGCAATTGGAATATCAGCTTCT
AGAAAACAATCTCCAATCTGAAAGAAGACAATCAGTACACCTGGCTATAACGGGTTAGCCTGCTAGGAGCCTGGTAACA
GCTGTGATTCTCGTAACGGCTTCTGTTAGTCAGTCAATTGGAAATGTCAGGAAGATTGTCATGCCAACAGTCAGTGA
GGGGATTCTGGTTAGGAATTAGCAGTACTATCAATCTGTTAGCAGTCTGGTGGTGTGAAAGGAAAGACAAGAAT
GAGTCTATTCTGAGTCTGCAATTCTGGAAGATACGCTAGGGTGGTAGCTGTTATCCTGATGGCATTGTTCTGATTAC

GGACTGGTATATCCTAGATCCTCTTTGTCATTCTTCTTATTCTTCAAAGCCCTCCACGTCTGGTCTACA
CTCAAGATTTCCTGGATGCTGCGAGAAGGTCTGATATCAAGCAAGTAAAGAGTGGCCTGGAGCGATGGACAATGTGG
CCAGCCTTAATCAGCTTAATCTCTGACTATGGATGCTTGAAAAAAATGCCATTGTCATGTTGCTAAAGAAATGGAA
CATATGGAAACTTGAAGAGTCTATTGAATTTCCTAAAGATTGTGGTTTCAAATATTACCATGAAATTGATGCTGA
CCTAGAAACTCACCAAACCCATAAGCAGAACGGAGTTATGAGCATCACATTAG

4170.8 (SEQ. ID. NO. 295)

ATGATTGAATACAAAATGAGCACTGCGCTACACAGAAAAGGATGTCAGAGAGATGTCACACTACAGATTGAGGATGGG
GAATTATGGTTTAGTAGGGCCTCTGGTCAGGTAAAGACGACCATGCTCAAGATGATTAACCGTCTTGGACCAG
TGAAATATTATATGGATGGAAAGCGCATCAAAGACTATGATGAGCGTGAACCTCGTCTTCACTGGTTATGTTTACAGG
CTATTGCTCTTCCAATCAGTGGCAGAAATTGTCATCCGTAATGAGGGTGGAGCAAGGAAGAAATT
ACGAAGAAAACAGAAAGCTTGGCTAAGCTGGTACCGTAGCCGAGTGGGCATCCCTACCTAGTGAATTATCTG
GTGGAGAACAGCAACGGGTCGGTATTGTCAGGATGATGGTCAGGCTTCAAGATTTCTCATGGATGAAACCTTTGGG
TTGGATGCTATTICGAGAAAACAGTGCAGGTTCTGACAAAAGAATTGATCAAAGAGTTGGGATGACAACGATTTGAA
CCCATGATACGGATGAAGCCTGAAGTGGCGACCGTATTGCTGCTTCAGGATGGAGAAATTGCCAGGTAGCGAATCC
CGAGACAATTAAAAGCGCCTGCAACAGACTTGTAGCAGACTTGTGAGGTAGTGTCTGACTAA

4171.1 (SEQ. ID. NO. 296)

ATGTCAGCAGTGGCTATTCAAGCTATGACCAAGGTTATGAAAGAAACCCACGGAAATCCTCTAGTATTGATGGTCATGGTCG
TCAAGCTGGCAAACCTTGCAGAACGCCCCTCAGGAACCTAGCCAGTTACTAAGGACAAAACCTCAACATATCTTTCACT
TCTGGTGGACTGAAGGAATAATACCATCATTGGCTACTGTCCTGTCACCAAGAACAGAAAACATATCATCACAA
CTGCCATCGACCATGCTGCTGAAACAATTGATTACTGGTCAACACACTTGGGTTGAAGCAACCATTATCCAGCCA
GAAAATCAAGAAATCACAGGCGAAATTCAAAAGGTTACGTCAGCATACGATTTGGTTTACCATGTTGTCATA
ATGAGACAGGAAACCTACTGCCATCGCTGAAATTGGCCTAAACTCAAGAACACCCCTGCTGCTTACATGTTGATGAGT
TCAGGCTATTGTAATCCCAATTCACTAGAAGAATTGGCATTGATTTCTCACTGCTCTGCCACAAATTCCATGGTC
CTAAGGGAAATCGGTTCTACGCATCTAGCATGGACTTGTGATTCTATCTACATGGCGAGACAGAACAGAAAACG
TGAGGAACGAAAATCTGCTGCCATTGAGGCATGGTGCAGCCCTAAAGAAGACCTAGAAAACAAGAACATTT
TCAACATGTAACAAATCTAGAAACTGCCCTTCTGGCAGAGCTGGAGGCATTGACTATTACCTGAAATAGAGGAAACATC
CTCCCTTATGTTCTCAATTGGATTCTGGTCAAGAAAATGACCTTCACTCCTGGTAGATTAGCTGGAATTCAATC
TCTACTGGCTAGCCTGACTGCAAGGCTTGTCCAATCAGGCTATGTTGAAAGCCATGATGGCGCAAAATTCAAGCAGCTT
GAAGGAATCCCTCGCATCAGTTGTCGCCACAAAATACCGTTGAAGACCTACAAACCTCGCAAAACCTTAAAGAAATT
ATCGGAGGTTAG

4172.1 (SEQ. ID. NO. 297)

ATGTTATTCAAAATTCTAAGGAAAAAAATAGAGCTAGGCTTATCTGTTATGCCAGCCCCGTCTGTTGAGTTTGGC
TTGGCTATTCTACTAGGCTCTCTCTTGTAGCTGCTTGTCAAGTGAAGGCTACGAGCGACTTATTTGATCATCTT
TCACTGCTCTCTGAGCTGTGACGGGCTCTCAACCCCTCCAGTAGCTCACACCTATAATATCTGGGTCAAATAATC
TGTGCTCTGATTCAAGATCGCTGGCTAGGGCTATGACCTTATTGGGTTTCTATATCCAGAGCAAGCAAAGCTTAG
TCTCGTAGCCGTGCAACTATTCAAGGATAGTTAGTTATGGAGAAACTCGATCTTGAGAAAGTTGCTATTCTATTCT
CACGACCTTTGGTGAAGAGCTTGGAGCTATTGCTTAGTTGCTTATCCCTCAACTGGCTGGGACGTGGTCTT
TAGTTCCATTCTTCTAGGATCTCAGCCTCTGTAATGCCGTTTGTATAATTAGGGAGCACCAGTTATGGCTTCTCAGAC
CGATTACTGGTCAATCTGTTGATTGAGCTGAGGCTTGTATTACAGGGGGCTTGGTTATGGTCTGGTTGATTGGCTGTCA
TGTAGGAAGAAAAGAAAAGGACGCTGCACTTCACTAGAAGCTTGTACTATTAGTACTAGGTTGTTATTGAA
CAGCAACTACTCTTCTGAGTGAACAATGCTGAAACGATTGCACTCCCTGTTGCCGATAAGGTTTAGTTAGCTT
TTCAACAGTGAATGCGAACAGCTGGCTTCTACGATAGATTACTCAGGCTATCCTGACTCTTGATTATATC
TTACAGATGTTCTAGGIGGGCCTGAGGAAACAGCTGGGGACTCAAGATTACGACATTTTGCTCTGGTCTT
ACGAAGTGAAGCTTCTAGGCTTGTGAGACAGCAACGATGCCGCGCGAACGGTCAAAAATCCTTCTAGT
GCTTATTATCTTCTGAGCTCTGATAGGATTGATTGCTAGGGATAACGCCAAGGCAATCCTCCCTTATCCAC
CTCGTATTGAAACCATTCACTAGCTTGTAGTACAGTTGGTGAACGGCAATCTGACTCCTGACCTTGGAAATTGGCTCTCAG
TGTATCATGCCACTTATGTTATGG
ACGAATTGGCTCCCTGACCTTGTAGCTGGCAGATTACCATCCAGAAAAGAAGATATGATTCACTATATGAAAGCA
GATATTAGTATTGGTAA

4172.2 (SEQ. ID. NO. 298)

ATGTCAGATCGTACGATTGAAATTGGGCTTGGGAAATTGGGAGCAGTGTCTAGCTGCCCTAGCCAAGCAGGATATGA
ATATTATCGTATTGATGACCAACGAGCGCATCAATCAGTTGAGCCAGTTGGCGCTGGAGTGTGGTACATCAC
AGATGAAGAATTATGGAGATCAGCAGGGATTGATACCTGCGATACCGCTGTAGTCAGCAGAGTGA
GGTCTGGGGTATGCACTGAGAGTTGGGGCTACGGACTGTTATTGCTAAGGCTAAAGCTAGACCCCTAAGGAAAGT
GAGTGTGCTTCTGAGCTGGCTTATCTGCCAGAGTATGAAATGGGAGCTCTAGCACAGACCATCTTCTCAA
TAGTGTGATGCTTCTGAGGATAAAATGTGCTATCTGGAGAGTGA
AAATTCTCTAGGCTTGGAGGAAAGTCAAGGCCAAGGCAATCCTCCCTTATCCAC
GTAATTAGACCTCCGTGGCAATACAATCTGAATATTGGGTTCCGAGAGCAGGAAAATCCCCATTGGATGTTGA
GGACCATGACCTCTGAAAGCAGATACCTATATTGGCAGTCATCAACAACCAAGTATTGGATACCTAGTAGCATG
ATTGTA

4172.3 (SEQ. ID. NO. 299)

ATGAAGTTATGCTATCGCAATTCTAGCTATAATGCAGCAGCCTATCTCATTACTGTCAGGAGTCGCTAGTGTG
TGAGCAAGTGGGATTGGTATTGATATCAATGACGGGCTCAGGATCAGACTCAGGAATCGCTGAGTGTGTTAGCTAGCAAGTAT
CCTAATATCGTATTGAGCCATCTACAGGAAAATAATGCCATGGCGGTGCGGTCATCGTGGCTGGTAGAGGCTCTGGC
GCTATTAAAGTAGGTTGACAGTGTGACTGGGATGCTCTGAGCTACTGAAATTCTGAAACCTTGCAGGAACCTGAG
ACGAAGGTCAGAGGGGAGTGTCTGAGCAATTGCTATGAAAGGAGGAGCAGTCTCGTAAGAAGAGTATGAGTT
ACGATTCTGACTCTGCTTGTGAGCTGACCAATTGCTATGAAAGGAGGAGCAGTCTCGTAAGAAGAGTATGAGTT
TCGCTGATTATCGGACAGATTGTTGCGTGTAGCCAGTTCTAA

4172.4 (SEQ. ID. NO. 300)

ATGAAATTCAATCCAAATCAAAGATACTCGTTGGTCTATTGCCGCTCACTGTCGGTGTGCCTCAGTTGTTGGCTAG
TGGCTTCTTGTCTAGTGGTCAGCCAAGTCTGTACGTGCCGATGGCTCAATCCAACCCCAAGTCAGTCTTACCTGAAG
AGACATCGGAACGAAAGAGGGTGAECTTACAGAAAACCAGGAGACACCGTTCACTCAAGCAACCTGAGGGCGTTA
CTGGAAATACGAATTCACTCCGACACCTACAGAAAAGACTGAAGTGAGCGAGGAACAAGCCCTAGTCTGGATACACT
TTTGAAGGAGATGAAGAACGCTTACAGGCTAACAGATGCTTAAAGAAAAGTGAAGTACAGCTGATGTGGA
TGGACACAAGCAAGTCAGCAGAAACTACTCTGAAACAAGTAAAGGGTGGAGTGAAGAAAATACAAAAGACAGCATCG
ATGTTCTGCTGCTTATCTGAAAAGCTGAAGGGAAAGGTCTTCACTGCCGTGTAACCAAGTAATTCTTATGAACTA
TTCGCTGGTATGGTATGTTAECTCGTCTATTACTAAAAGCTCGATAATGCTCTTGGTCTGACAATGGTACTGCTAAAAA
TCCGCTTACCTCTTGTAAAGGATAACAAAAGGAAATACTCTATGAAAGTAACTTAAATGGCAATACTGTTGGTAAAC
AAGGTCAAGCTTAAAGTCAACTCGCCTAATGGTACTCAAACCTATAAAGCTACTGTTAAAGTTACGGAAATAAAGA
CGGTAAGCTGACTTGAATCTAGTGTCTACTAAAATGAGCATCACATCAATGCTTAAAGGATAGTTGCTTAAAGAAAACAGTT
CAAAAGCCGTTGAGACAACGTTAAAGAGCTATCGATGTTCCAGCAGCCTACCTAGAAAAGCCAAGGGTGAAGGGTCCA
TTCACAGCAGGTGTCAACCATGTGATTCATACGAACCTCTCGCAGGTGATGGCATGTTGACTCGTCTTGTCAAGGCATC
TGACAAGGACCCATGGTCAGATAACGGCAGCCTAAAACCCAGCCCTATCTCCACTAGGCGAAAAGCTGAAGACCAAAGG
TCAATACTCTATCAAGTAGCCTGGACGGAAATGAGCTGGCAAGGAAAACAAGCGCTATTGACCAGTCCGAGCAAAT
GGTACTCAAACCTACAGCGTACAGCTCAATGTCATGGTAAACAAAGCGTAAACAGACTTGGATTAGTTGCTTAAAGAAAACAGTT
AAAAAGTCACTTAAACATAAAAGCTTAAATTCTAAAGAAAAGCTTAAAGGCGTTGAGACAACGTTAAAGACAGTAT
CGATGTTCCAGCAGCCTACCTAGAAAAGCCAAGGGTGAAGGTCTTACAGCAGGTGTCACCATGTTGATTCCATGAA
CTCTCGCAGGTGATGGTATGTTGACTCGTCTTGTCAAGGCATCTGACAAGGACCATGGTCAAGATAACGGTGACGCTAA
AAACCCAGCCCTATCTCACTAGGTGAAAACGTGAAGACCAAAGGTCAATACTCTATCAATTAGCCTGGACGGAATGTA
GCTGGCAAGAAAACAAGCGCTATTGACCAACTCCGAGCAGCGTACTCAAAACTACCGCCTACAGTCATGCTATG
GTAACAAAGACGGTAAACCAAGACTTGGACAAACATCGTAGCAACTAAAAAGTCACTTAAACATAAAACGGTTAATTCTAA
AGAAACAGTCAAAAGCCGTTGAGACAACGTTAAAGACAGTATCGATGTTCCAGCAGCCTACCTAG

4172.5 (SEQ. ID. NO. 301)

ATGAAACTAAAAGTTATTTGGTGGATAATTTCACCCCTTAAACCATTGGTTGTTGGCTGTTCAAAAAA
ATGCTGATTGCGAAAGGGAGATTACTTTGCTTGGGATGACCATGTTGCCAGCCTGTCGGTGTGGATTAGTCTCTT
TCTCTTATTGCGACTTACGTCGTTGGCAACTCAAGGAGCATGCCAAGCGGGTAGCGGCAAGGATTTCCTCAAAATT
TGGAGGTCAAGGCTGTAGAATTTCAGCAATTAGGGCAACCTTTAATGAGATGTCCTGATTTGAGGTAAAGCTTGT
TCCTTGAAGAAAGCGAACGAGAAAAGGGCTGATGGCCAGTTGCGATGTTGCGCATGATATTAAAGACTCTATCCTGATCC
AAAGCAGCGTAGAAGGGATTGGATGGATTATCAAGGAGTCGGAGCAAGCTCATTATCTAGCAACCATTGGACGCCAGA
CGGAGAGGCTCAATAAAACTGTTGAGGAGTTGACCTTGGGCTAAACACAGCTAGAAATCAGGTGAAACTACCAAGTAA
AGACAGTATTCTGGACAGCTCTTAAATTGAGTCATGAGTGAATTCTAGTTGAGTGGAGAGGAGAGAGATGTC
ACTTGCAGGTATTCAGGAGTCTGCCCAGTTGGAGGATTATGCTAAGCTTCTGATCTGGTGAATCTGGTCGATAAC
GCTTTAAATATTCTGCTCAGGAACCAAGCTGGAAGTGGCTAAGCTGGAGAAGGACCAAGCTTCAATCAGTGTGACCG
ATGAAGGGCAGGTATTGCCCCAGAGGATTGGAAAATTTCAACGCCATTCTGTCGAAACTTCGCGTAACATGAA
GACAGGTGGTATGGATTAGGACTTGCAGTGGCGTGAATTGGCCATCAATTGGTGGGGAAATCACAGTCAGCAGCCAG
TACGGTCTAGGAAGTACCTTACCCCTGTTCAACCTCTGTTAGTGAAGAAATAAGCCTAA

4172.6 (SEQ. ID. NO. 302)

ATGTTGGTCAACATGGCTTACGAATAGCCTGAAAGACTTCTGGATTTCCTGCTGAATATAGGTCCACAATT
GGCGTTTTGCGAGTGCCTCGCTGTTCCAGTCGGTGGAGCAGGGTACTGGAATCACCCTGCTGAGTTCAATATGATTTC
ACGAGATATTCTGCATTGGGATGACTCACTTGGGACAAATCAAGTGGCTATCAAGACTGCGATTGACCTTGAGTTGCTG
GTCATGCACTTAATCATCACTTGCATTGACAGACTGGCTCATTGACAGACTGGCTCACGCCAATCAAATAACGATAGAAATCGACAGCAGA
TAGTACATGGCTTGCACCTGCTGAAGGGCGTAAAGACAAAGAGATTATGACATAAAAGTATGTCAGGAGCTTCAATCAGTGTGACCG
GGCTAGCACCAACAAATCTGTCGATAATCAGCAGTCATGGTACTGGCCTTGGAGAAATTCCGACCTGGTC
CCAGCAAAATCTGCCAACAGGAAGACTGA

4174.1 (SEQ. ID. NO. 303)

ATGGAACATTTAGCACTTATTTCAACCTATGGAGGAGCTTCTCGCTGCACTGGAAATTGATTGGGGTTGGATTAAG
CGGTATGGGCTGCTTATGGAGTGGTAAGGCTGGCAATCTGCCAGCTTACTGAAAGAACAGCCTGAAAAGTTGCC
TCAGCTTGTATTGCAATTATTGCCGAAACAAGGATTATGGTTATTGGAAATTITAATTGGTGAATTAAACT
CCAGAACTTCTTTAGAAAAGGGCTGCTTATTCTGCTAGCTTCAATTGCTATTGAGGACTATTCTAGCTAAGCAT
CAAGGAAATGTAGCAGTAGCGGAATGCAAATTCTGGCTAAAGACCAAAGAGATTATCATGAAGGGAGCAATTAGCTG
ATGGTAGAAACCTATGCAATTCTGTTTGTGCTATCTGATTGACCTTCTGGTATAAA

4175.2 (SEQ. ID. NO. 304)

ATGTTAAAATCAGAAAACAATCACGTTATCAAATGTTAAATGAAGAATTGCTCTTCTATTGGAGGGCAAACCAATGTTT
TGCTAATCTTCAACCCAGTCAGTCCTCATAAAATCACGTTCTCTAATACCGTATTGCAAGGCTTATTGTCAGTGGAA
AGGAATTGGTTTAGGCCCTTCAAGGAGGTTCTCTGATCCGATCCGTTATGCACTAGGCAAGGGTGTGAGGGCAGCT
CACTTCAAGGAAACTGTTATTGAGGATGTGACGACCTATCTCAACTATATTCTGCTAGTCTAGCTAAAGTGAAT
TGTGGTGGCGATGATGAAGAATGGTCAGTTGGAGGTTCTGGATCTGAGTCTGAGAGATTGAGGATACGATGCTATGG
ATCGAGATTATTGGAACAATTGTCGCTATTGGCTGAAAGACAGCATGGACTTACGATGTTGAGGAAAATCTAA

4175.3 (SEQ. ID. NO. 305)

ATGTCAGTATTAGAGATCAAAGATCTTCACGTTGAGATTGAAGGGAAAAGAAATTAAAAGGGGTTAACCTGACCCCTGAAAA
CAGGAGAAATTGCCGCTATCATGGGACCAAATGGTACAGGTAATCGACTCTTCTGCCGCTATCATGGGAAATCCAACCA
TGAAGTAACAAAGGTGAAGTTTGTGATGGCTAACATCCTTGAGTTGAAGTGGATGAGCGTGCAGCTATGGACTT
TTCCCTGCTATGCAATACCCATCAGAAATCCCTGAAATTACCAATGCTGAGTTCTCGTGCCTATGAAATGCGGGTAAAGA
AGATGATGAGAAGATTCTAGTCTGAGTTTACTAAGCTAGATGAAAATGGAATTGCTCAACATGAAAGAAGAAGAATG
GCAGAGCGTTACCTCAACGAAGGGCTCTGGTGGTGAAGAAAACGCAATGAAATTCTCAACTTTGATGTTGGAGGCAA
CATTGCTTTGGACGAGATTGACTCAGGTCTGATATTGACGCTTAAAGGTGTCATGCCATCGT
GGTGAAGGTTTGGTGTATGATCATCACTACCAACGCTTITGAACATATCACACCTGATGTTGACACGTGATGAT

GGAAGGTCGTGTTGTCCTTCTGGTGGTCCAGAATTGGCTGCGCTTGGAACGTGAAGGATACGAAAATTAGCTGAAGAA
CTTGGCTACGACTACAAGGAAGAATTGTAAGA

4174.4 (SEQ. ID. NO. 306)

ATGCCCTACAAAAGACAAAGGAGTTTCAATGGCACTTCTAAACTAGATAGCCTTATATGGCAGTGGTAGCAGACCATT
CGAAAAATCCACATCACCAAGGGAAAGTTAGAAGATGCTGAGCAAATCAGTCTCAACAATCCGACTTGTGGGATGTCATCA
ACCTCTCTGTCAGTTGATGCAGAGGACGTTGGAAAGATATTGCTTCTAAATTCAAGGATGCACGATTCAACTGCTTCT
GCTAGTATGATGACAGATGCCGTTAGGAAAAACCAAGAAATTAGAACACTGGCAGACTTTCTGAAATGGTTC
AAGGGAAAAAGATGAGCGTCAAGACCAACTGGAGACCCGGCATCTGTCAAGGTGTTGCAAATTCCCTCAAAGAACATCA
AGTGTGCAACCTAGTGGAAATGCCCTAAGAAAACAATTGAAAATCAAGAAAACAGTAA

4175.5 (SEQ. ID. NO. 307)

ATGAAAATTCAAGACCTATTGAGAAAAGATGTCATGTTGCTAGATTGCAAGGCAACTGAAAAAACAGCTGTCATCGACGAGA
TGATTTAAAATTGACAGACCACGGTTATGTAACAGATTGAAACATTAAAGAAGGAATTGGCGCGTGAAGCTTGTAC
TTCTACTGGTTGGGTGATGGAATCGCAATGCCCTCACAGCAAAACGCTGCTGCAAAGAAGCGACAGTCTATTGCTAAG
TCAAATAAGGGTGTGACTACGAGAGCTGGATGGACAAGCAACTGACCTCTTCTCATGATTGCAAGCTCCAGAAGGTGCCA
ATGATACTCACTGGCAGGCCCTGGCAGAATTGCTCAACTTGATGAAAGACGGTTTGCAAGACAAACTCGTCAAGCAC
ATCTGAGACCAAGTTGACCAAGCTTCAAGGCTGCAAAGGAAACTTGCTTCAAGCACCTGCTAATGACTCT
GGTACTTTATCGTAGCTTACAGCTGTCACAGGTATTGCCCACACTTACATGCCCAAGAAGCCTTCAAAAGTAG
CTGCTGAAATGGGGTTGGTATCAAGGTCGAAACCAACGGTCTAGGGTGTGAAATCAACTAATGCAAGAGATATCCG
TAAGGCTAAAGCTATTATCATTGCAAGACAAGGCCGTGAAATGGATGATTGATGAAAACCAATTGATCAATCGTCCA
GTTGCTGACGGTATCCGTAAAGCAGAAGAGCTAATTAACTTGGCTTTCAGGAGATACTGAAAGTCTACCGTGCCGCTAATG
GTGCCAAGCTCAACGCCCTCAACGAAAACAAAGCCTTGGTGTGCCCCTGTCACAAACACTTGATGAGTGGTGTATCTCA
AAATGTTACCTCGTTATCGGTGGTGTATCATGATTGCCCCCTGCCCTTGTGATTGACGGTGTGTTGGTGTGTTCAAATGAAA
CCTTGGCAATCTGGTCTTACCATGAGTTAGCTTCTATGTCATGAAAATTGGTGGAGCTGCCCTTGGTGTGTTCAAATGCTTCA
CTTGGGGTTATGTTGCCACTCTATTGCTGAAAACCAGGTTGGTAGCAGGTTCTGTCGGCTGGTGTGTTGCAATTGCAAAAGAAG
GTTTGCCTTGGTAAAATTCCATTGCCAGGTGGTGAAGCAACTTCAACTCTGCAAGGTGTCATCTGGTTCCCTAGGTG
CCCTTGGTGGGATTATCGCAGGTGCCCTGGTCTGCCATCAAGAAATACGTTAAAGTCTCGTCACTCGAAGGTGCT
AAATCAATCTTCTTCACTTCCACTTGGCAACAAATCTGACAGGTTGTTGACTAGCTGTCAGGTTGCAATATCCCAATGGCTCAAT
CAACACTGCTAATGATGACTTCTCAGGCGCTTGGAGGAGTTGCTGTCAGGTTGCTTGTGTCAGGTTGCAACTGTTCTCAGGTG
CTGTTGACATGGTGGACCAGTAAAGCAGCTTATGCTTGGTAGCAGGTTACGCTGCAAGCAACTGTTCTCAGGTG
TCTGTAGCCATGGCAGCAGTTATGGCTGGAGGAATGCTGCCACACTGCAATCTTGTGCAACTCTTCTTCAAAGATAA
ATTACTAAGGAAGAACGTAACCTGTTGACAAACATCATGATGGCTTGTGTCAGGTTGCAACTGAGGGAGCATTGATTG
GTGCCGCTGACCCAGCTGTCGCGATTCCAAGGCTTCATCCTTGGTTCAGCAGTAGCAGGTTGACTCCTGGTCTTACTGGTATC
AAACTCATGGGCCAACACGGAGGAACTCTGTTATGCCCTTACTTCATGTCATGCTCCCTTACCTCGTTCTGCTTGTGTTGAGGA
GCAATGTAAGTGGTGGTTATGTTACCTACGAAACCAAGCATAA

4175.6 (SEQ. ID. NO. 308)

ATGGCAAACAAGAAATACAAGTACAACAAGACGGAGACCGTCTAAAGCAGAACTGGAAAGAAAAGAAGCGATTCAACGAAT
GTGATTTCTGTTAGGAATTGCGATTGATTTATTGATTTGCAAGCCTCAAAATTAGGGCTGAGGTATAACCCTTATAATTAAAT
TCGCTTGTCTAGTGGTAGCCTAGCTTATCTGGCGATATTGCCCTATTAAATCTATCTCTTCTTCAAGTGGATACGAAAACA
GGAAGGACTTTATCTGGCTTTCACCATATTGCTGGCTACTCTGATTTGAGGCCACTTGGTTGAAATATGGTT
GGACAAGTCCGTTCTAAAGGGACCATGGCTCAGGTTGACAGATCTGACTGGTTGCAACGACTAGCTTGTGAGGG
GGCTTGATCCGGTGTCTTATATTCAACAGCCCTTCTTCAAAATATCGGAAACTTACTTATTGGTCTATCTTGATT
TAGGGTTCTCTTCACTGAGCCCTTGGTCTTACGATTGCTGAAATTTCAGTAGAGGCTTGGCCTTCAAAATGGGGAA
GGGCAGGCGTCAAGAAAAGGAGAACGGTCTGCAAAAGAAGAAAAGCTGCCAAAAGGGTGTGAAAGAGGCTAGATT
AGAACAAAGAAGAGACTGAAAAGCCTACTCGATTGCTCTGTTGATATGAAACGGGTGAAATTCTGACAGAGGAAGC
TGGTCAAATCTCCACCTATTCCAGAAGAAAAGTGGGTGAAACCAGAAAATCATCTGCCCTCAAGCTGAACTTAAATCCCT
GAACAGGAAGATGACTCAGATGACGAAGATGTTAGCTCAGGTCATTTCAGGCTTCAAGAAGCCCTGAAATCAAACCTTCAAGCT
TACAACCTTCTGACCAGATAAAACAAAAGATCAGTCTAAAGAGAAGAAAATTGTCAGAGAAAATATCAAACATCTTAAAG
CAACCTTCTGCTTGGTATTAAAGGTAACAGTGAACGGCCGAAATTGGGCCATCTGACCAAGTATGAAAGTCAAGCC
GGCTGTTGGTGTAGGGTCAACCGCATTCCAACTATCGGAAATTGAAAGTCTGCTCTAGCCTTGGCTGCCAAAGATGTCGGATTG
AAGCACCAATCCCTGGGAAATCCCTAATCGGAAATTGAAAGTCTGCCAACTCCGATATTGCCACTGTTATCTTCCGAGAACTATG
GGAACAAATGCCAAACGAAAGCAGAAAATTCTGGAAATTCCCTTAGGGAAAGGCTGTTAATGGAACCGCAAGAGCTTGTGAC
CTTCTAAAATGCCCTTGGTCACTGCTAGTTGCAAGGTTCAACGGGTTAGGGAAAGTCAGTAGCAGTTAACGGCATTATTGCTAGCAT
TCTCATGAGGCCAGACAGATCAAGTTAAATTATGATGTCGATCTGGCAGTCTGGAGTTATCTGTTACAAATGATATTG
CCCACCTTGTGATTCTGCTGACCAATCCACCGCAAAGCCAGCAGAAGGCTCTGCAAAAGGTTGTTGATGAAATGAAAACCG
TTATGAACTCTTGTCAAGGTGGAGTTGCGAATATTGCAAGGTTTAATGCCAAAGGTTAGAAGAGTTCAATTCCAGTCTGAGT
ACAAGCAAATTCCGCTACCATTGCTGTTGAGTTGCTGAGTTGGCTGACCTCATGATGGTGGCCAGAAGGAAGTGG
AGATGTCATCATCGTCTGGCAGAAGGCGCTGCTGCAAGGTATCCACATGATTCTGCAACTCAGCGTCCATCTGTTGATG
TCATCTGTTGTTGATTAAGGCAATGTTCTCTGCTGAGCTTGGGTTCTACGGAACAGACTCCCGTACGATTGTT
ATGAAAATGGAGCAAAAACCTTCTGGTGTAGGGAGACATGCTTITAAACCGATTGATGAAATCATCCAGTCTGCTCCA
AGGCTCTTATCTGGATGACGATGTTGAGGGCATTGTAACCTCATCAAGACTCAGGCGAGATGCAAGACTACGATGAGAGT
TTGATCCAGGTGAGGTTCTGAAAATGAAGGAGAATTTCGGATGGAGATGCTGGTGTGATCCGTTTGAAGAAGCTA
AGCTTGGTTATCGAAACACAGAAAGCCAGTGCCTATGATTGAGCTTACGCGTCTTACGTTGATTTAACCCTGCGACCCGT
CTCATGGAAGAAGTGGAGATAGCAGGTGTCATCGTCCAGCTGAAGGTACCAAACCTCGAAAAGTGTACAACAATAA

4176.1 (SEQ. ID. NO. 309)

ATGAGTTTAAAAATATAAATCGATAAATCCCAAGTCAAACTTGGTATGCGAACCTTAAACAGGTATTGCTGTTT
TCTAGTTCTCTGATTGGCTTGGCTGGAAAGGTCTCAATTGGTGTGTTGACAGCCCTTGGCTGAGGGAGAG
TTTGATGAGGAGTGTCTGGCTATTGGGACTTCGCTGTTAGGAAATAGTATCGGTGACTCTATGCCCTGGCTTCTTCTTATT
AAATACCTTCTGCCACGAAGCCCTTGGGTGACCTTGGTAGTTGCTTCAATCTGACCATGTTAACCTTATGACAATGTAG
CCATGAATACAAAGCAGGGTTATTGGTGTGAGGAGACATGCTTCAATCTGACCATGTTAACCTTATGACAATGTAG
TTTGTACGTGTTGCGTGTATTAGAACGTTATGGGAGTTTGTGCAATTATCGTAATTACGATATTGATCGTATTG
TCTCTTCTAGAGAAAAAGAAAAATAA

4178.2 (SEQ. ID. NO. 310)

ATGAATAAACAGAACACCGCCACCAACTTATACGGCCTTATCACAAAAAAACAAGATTACACAGGCTGAGTTGCAAG
CCCTTCTGCTGAGAACGACATTCAAGTAACCCAGGCAACCCCTCTCACGCACATAAAATATGAACCTATCAAAGTCG
CGAAGAAGATAGCGCTTATTATGTTCTAACATGGTCCATCTCAAATGGGAAAAACGCTCGAACTCTACATGGAAGAC
GCCCTTGTCTGGATGCCCCAGTCAACACCAAGTCTACTAAAACCTTCTGGACTGGCTAACCTTGGTCTATTCTATT
GATACTTGAGCTTCCCTGACGCTATCGCTACCCCTTGTTGAATGATGTCCTATCATCTGTGAAGATGAGATACTGCT
CAAAAGTCTTGAAGAAGTGAACCACTGCCCCACCATTTCCTTGAAGAATAA

4179.1 (SEQ. ID. NO. 311)

ATGAAAGTATAAAATTAAATGCTCATCTTACATGGGAATTCTGTCCTGAAATTATTTCCCACCTAACTGGAACCTA
TGTGCGCGTGTCTGGACCGAACGACTGACTATGGTACTTCAACTCAGTCGACACTATTTGTCATTTCCTGGCTTGCAC
TTATGGTGTCTATAACTACGGTTAAGGGCTATCGTAATGTCAGGATAACAAAAAGATCTAACAGAACCTTCTAGTC
TTTTTATTTGTCATCGCTGTACGATTTGACCACTGCTGCTATATCTAGCCTATCCTCTTACTGATAATCCAAT
CGTCAAAAGGTCTACCTTGTATGGGATTCAACTCATTGCCAGATTTCATCGAATGGGTAATGAAGCTGAAA
ATTACAGTTCTCTTACAAACTGCCATCCGTATCTGATGCTGGTCTCTATTTCATTTGTTAAAATGAACACG
ATATTGTTGCTATACACTTGTGAGTTATCGACGCTGATTAACCTACTCTGATTAGTTTGGATTAAGAGACATCA
AACTGTTAAATTACCTAAAGTATTAAACACTCTTCCCTCTGACGCCATGTTAGCTTGGCAATGCCATATG
TCTTCACTTTTAGATGCCCTTCTCGTTAACACGGGATTGATGTCACGTTAGTTACTATACCATAGCTCAGGAATTG
TGACCGTTAGCTGGGGTTGTAACAGGTGCAATTGGAGTGTGAGTGTGCTCGTCTAGTTACTATCTGGGAAAGGAGACAA
AGAAGCCTATGTTCTGGTAATAGAGGTAGTGAATCTTAACTCTTATCATTCACTGAGTTGGACTCATGTTT
AGGACCAAATGCCATCTACTTACGGTAGTGGAAAATATCGGAGGGCAGCTTGCACCTCTCTCCGTTTGTACGA
TTACCTGGCCTTAGATACCATTCTGGTCCAAATTCTTACCATGGGCTATGAAAACGCTACAGTCTATACAGTCT
TTGCTGGGCTACTCAATTGGGTTGTAATGTCCTTTCATCGCTGCTCTGAATACTTACTGACAACTA
TGCTATCAGAGACTCTCTACTGTTCTATATCATTTCATCAGAAAACAACCATCCACTGGGACATATCTTAGCT
ATACTGTTGATACTCTCTTCACTTCTGTTAGCAATTATTCCTGATTAATTCTGTATCTGTAGATATGGTCAT
TAATTGCCATTGGTAA
ATACTGGTTGATTGTCCTGCTATCAGCTATCTTATATTAGTCTACTTGTCTCACAAAGATAGCATTTCTATGAATT
AAACCATGTCTAGCCTAAAAAATAAATTAAAATCATAG

4179.2 (SEQ. ID. NO. 312)

ATGAAACAACAAACGGTTGAGATGCCAACAAATTGAATTAGAAATTGGATTATATTGATACTCTGTAAGCACA
ATATCAACTATTTAAACTACGGTACTCTGATTGGGGCGGTTGACATGAGGGCTTATCCCTGGGACGACGATATTGAT
CTGTCATGCCAGAGAACACTACCAACGATTATTAACATTTCAAAAGGAAAAAGCAAGTATAAGCTCTATCCCTAG
AAACTGATAAGAAACTACTTAAACAACTTATCAAGATAACCGACAGTACGACTAAAATTATTGATACTCGAAATACAAAAC
CTATGAGTCTGGTATCTTATCGATATTCCCTATAGATCGTTGATGATCCTAACGGTATTGATACTTGTATAAAACTGGA
AAGCTTCAAACTGCTGTTCACTGAAACATAAAATATTGCTATAAGGATAGCCTTAAAGGATTGGATACGAACAGCCT
TCTGGTTACTCTCGACCCGGTTCTCTCGTTATTGCAAAATGGAGAAAGAAGGAAATTCAAAATATAGTCGTTGAAAAT
GGCAATATGGTTTATCCCTCAAAATTAAAGGAAAGTCTCCCAAGTGGTACCTTGTATAAAACAATGATT
ACCCITTGAGAATTAAAGCCTCTGACCTGAAATTGATACTATTGACACAAATTATGGAGATTATGACCCCTAC
CACCAGAAGAAAACGCTCTACAGTCATGAATTCACTGCTATAATTGGAGGATTAG

4179.3 (SEQ. ID. NO. 313)

ATGATAAAAATCAATCTAACCATCACACAAAACAAAGTTACGAGATCTGATCTGACCTAACCATGACCATCCAAG
ACGGGGAAAAGGTTGCTATTATGGTAAGAAGGAAATGCCAAATCAACCTTACTTAAATTAAATGGGGAGCTTGT
TGATTTCATCTACAGGAAACATCCAATCTGACTATCAGTCACTGGCTACATTCTCAAAAAGTCCCTGAGGACCTAAA
AAGAAAACCTTACAGACTACTTCTTAAAGTATTGACTACAGTACAGTATCTCTATCGTTGGGGAGGAATTG
TTGATAGCAATCGTTGCAAGTGAAGAGATTGGCAATCTATCAGGGGGCAAGGTTGAAATTAGCTTATCCATG
AGTTAGCCAACCCCTTGTGAGATTCTTAAAGTGAACCTTCAATGACCTAGAGACTGAGCTTGTGATGGCTAAAAGG
CAGATTCAAAAGACGAGGAAACCGTTATTCATTTCCATGATGAGACTTCTCTGAAACGGCAGACACTATTG
CTTGCAGACTGTCACACCGTAAAGAAGCGGAAACGCTAGTAGAGCATTTGACTATGATACTGAGCAGAGAAA
GGCTAATTGCCAACAAAGTCAGCAAGCTGTAACAACCAAAGGCTACGATAAAACCATGGAAAACATGGAGAGT
TAAGAAAATGTAGAAACTGCGCTTCAGCTACCAAAGATAGTACTGCCGTCGCTATTGGTAAAAAGATGAAAATG
CTCTCACAAAGAAAACGCTACGAAAAGGAGCTAGTCCATGACTCAAAGCCACTTGAAGAGGAACAAATCCA
ACTTTCTTTCAGACATCCAACCATTCAGCTTCAAAAGTCTTACTGCAACTGGAAAAGAAAATTG
GTTCAAAACTACAACAACTGTCCTGGGCAAGAAAATGGTATTCTG
GCCAAATGGTTGGGAAATCAACTCTGTTAGCCAAGTTACAGAGACTCTGAATGATAAAAGAGAGATTCACTGGTTT
ATGCCACAAGATTACCAAAAAACTGCAATTGGATTATCCCAATAGCCTATCTCAGTAAAACTGGGAAAAGAGGAA
CTACAGAAAATCCAATCTCACCTAGCTAGTCTCAATTCACTGTTATCCAGAAATGCACTAACATTGCTCCTTATCTGG
ACACAGGGAAAACCTCGTCTTGTGATTAGTCTCGCAGAACCCAAACCTTCTCTGCTGGATGAAACCCACAGAAACTTT
CTCCCACTCTCAACCCAAAATCAGAAAACCTTGTACCTATCCAGGGGTCTCATCACTGTTCTGACGACCGAGTTG
TAAAAGAAGTCTGTCGATCATCTCGCATGACAGAACACGGTTGAAGCTAGTTAATTAGAAGATTATAA

4179.4 (SEQ. ID. NO. 314)

ATGAAACCAAAACATTTACAACCTTGTGCCAGCAGAACCTTCCACTTCCGACCGAGCAAAAGAACAAATTGAACGTT
ATTITGAGCTTGGTCAGTGGAAATGAGAAAGATTAATTGAGGGGATTACGGACAAGGAAGTTATCTCAAACACATT
TTACGATTGCAATTGCAACCCATTCTCAAGGTTGATTCCTAACGAAACTATCAAACCTTCTGATATCGGGGCTGGGGCAGGAT
TCTCTAGTCTACCAATGAAAATTCTCTATCCGGAGTTAGATGACCTATTGATTCTACTCAATAAGCGCATCAACTTCTA
CAACTCTGGCTCAAGAACGTTGAAACGGAGTTCTACCCACGGACGTGCCAGGTTGAAAGATTGCTTCAAGAACAGAAC
TCCGTGCTCAATATGATTGAAACAGCTCGTGCCTGGCCGTTGAGGATCTGAGGTCTATCTGAATTGACTATTCC
TTGGTGGCAACTATTAGCACTCAAGGCTAGCAATTGCGCTGAGGAATTATTAGAAGCTAAGAATGCCCTCAATCTC
AGTAAGGTCGAAGACAATCTCAGtACGCCCTACCGAATAGAGATCCGGCTATACACAGTGGTAGAAAAGAAAA
CACCAAAATAATCCACGTAAGGCTGGTATGCCAAATAAACGCCACTTAA

4179.6 (SEQ. ID. NO. 315)

ATGAGTATTAAACTAATTGCCGTTGATATCGACGGAACCCCTGTCAACAGCCAAAGGAAATCACTCCTGAAGTTTTCTGC
CATCCAAGATGCCAAGAAGCTGGTCAAAGTCGTGATTGCACTGCCGCCATCGCAGCGTTGCCAAACTCTAGAC
GACTTGCAGTTGAGAGACGAGGGGACTATGTGTAACCTCAACGGGCCCTGTCCAAGAAACTGCTACAGGACATGAGA
TTATCAGCGAATCCTTGACTTATGAGGATTATCTAGATATGGAATTCCCTCAGTCGAAGCTCGTGTCCACATGCATGCCATT
ACCAAGGAGCGGTATCTAATCGCAATCGAATATCGGAAAATACACTGTACACCAACTCACCTCGTCAACCATGCCATT
TCTACCGTACCCCTGAGAAATGGCTGGCAAGAAAATGTTAAATGTATGTTATCGATGAGGAGCAAAATTCGATGCTGC
GATTGAAAAAATTCCAGCAGAATTTCAGCGCTACTCCATCAACAAATCTGCTCCTTCTACCTCGAACCTCTAAAAGA
ATGTAGACAAGGGTCAGCCATTACTCACTTGCTGAAAACCTCGGATTGACCAAGAGATGAAACCATGGCAATCGTGTGA
AGAAAATGACCCTGCCCAGCTGGAAAGTCGTTGAAACCCGTTGATGGAAAATGGAATCCAGAAATCAAAAAATCGC
CAAATACATACCAAAACAAATGACGAATCCGGCTTGCCCAGCCATGCCATCCGAACATGGTACTGTAA

4179.7 (SEQ_ID_NO. 316)

ATGACTTGGATTATTCTGGAGTTATCGCTCTTATTGTTATTGTGATTGTTAGCTATAACGGTTGGTAAAATCGTATG
CAAACCAAGGAGGCTGGAGTCAGATTGATGTTCACTGAAACGTCGAATGACCTTGTCCAACCTGATTGAGAGCTGAA
AAGGTTATGCCAATATGAAGGTTACCCCTGAAAAGGTGGCAGAACTACGTAACCAAGTGGCGCAGCGACTTCACCAGC
AGAACGCTATGAAAGCCAGTGTGCCCTCACTCGTCAGGTTCACTGAGTTGAGGAGCTATCCAGATTGAAA
GCTAGTGTCAACTTGTAAATTGCAAGAGGAGTTGACAAACAGAAAATTCCTACTCTCGTCAACTCTATAACAA
GTGTTGTCAGCAACTACAATGTAAAATAGAAACTTCCCAGCAATTATCGTGGAAATGTTGATTAAAGCGGCAGA
TTCCCTCAACACCTGAAGAGGAAAAGTCGGTCTAAAGTGATTGTTAGCGGTTAGGTGACTAA

4179.8 (SEQ_ID_NO. 317)

ATGTTGTTGATCAAATTGCAAGCAATAACGAAAACCTGGATTITGTTGCTGGTATTITCTACTCTTAGCTTGTGGT
TATCGGGTTGGTTATCTTTATAAGATCTGGACTTGGTGGTTATTGCACTGATTATCGCTTATCTACGCTTGTCT
ATGATTTTCATCGACAGAGATTGTCATGTCCATGAATGGAGCGCTGAGGTGGATGAGCAACGGCACAGACCTCTACC
ATGTAGTGGAAAGATATGGCTCTGGCTCAGATTCTATGCCCCGTGTTICATCATTGATGATCCAGCCTTAAATGCCCTTG
CGACAGGTTCAATCTCAAATGCCCTGTTGCTCGACTCAGGTCTACTAGCTATCATGAATCGTAAGAAACTAGAAC
TGTATGGGACATGAAGTCAGTCATTGTAATTGATATCGTATTGCACTATTGAGTTGCTAGTGTCTGAGTGTATCAC
CATGCTTCTAGTATGGCAGGCTGTATGATGTTGG
GGTCTGAAATCATTATGCTAGTGGTCTCCCTACTAGCTATTGCTACTGGCACCTCTCGTCAACCTGTTGAGCTCGTATT
TCTCGTCAAGGGAAATTCTGGCAGATGCATCTAGTGTGAGCTGACTCGCAATCCCCAGGGAAATGATTAATGCCCTAGATA
AGTGGACAATAGCAAACCTATGAGTCGCCACGTCGATGATGCTAGCAGTGGCTTATATCAATGATCCTAAGAAAGGTGG
GGGGTCCAAAACCTTTTACCCACCCACCTATCTCAGAACGGATTGAACTTAAACAGATGTAA

4179.9 (SEQ_ID_NO. 318)

ATGAAATTAAATATTCAAGAAATTGTAAGCAGTCTGAAGGTTGAACCAAACGTTAGATTAGTTGATGACCTGC
GTGCACGTAATCAAGAAATTITAGATGAAAAGATATCCTGCACTGGGAAAGTACAATATGAAGACCTATGATTCTT
AGATTATCAACTATCTTACCAATTGTTCTGCTCGAGTCAGATGGAGCCAGTTGAGTTAGTTGAATCTTATCCAGTCAC
GGAAAGTTTCTGGAAGGCAGCAACTAACAGCTAGATCAAGAAAGTTTGTAGTGTACTTGGCTTGCACCATGAAAATGGG
GAGCTTGACCTTGTGAGAGTGTATCAGACAATATCCTGCTAAACATCTCAAGGTCTTGACGGCTGAAGAAGAGCTG
GTCAAGGATTATCTCAGGAAATGACTGGCAACATCGACAGAGGAAATACCAAGCTCAAAAGCAGTAAAGAAAGAAG
AAAACAGTCCTTGTGGCTTACAAGGACTATTGACGGAGATGAATAA

4179.12 (SEQ_ID_NO. 319)

ATGGAGTTATTGAAATCACAACATGAAATCTATAAGTTAAAAAAATCAGGTTGACCAATCAACAGATTGAAAG
TGCTAGAATACCGTGAAATGTTGATCAGGAGCTTGTGGGTGATATTGAGATATCTCAGGTTGCCGTAAATCCAGCGTT
TTATGGAACGTTATTCAGATAGACGATGCCATTGTCGAAAGAGTTCAAAATTCATCTTCTATTAGATGAC
TGTATCCTGGGATTGAGGAAATATGATGCGCTGTACTTTATTACAAGGGAAATCTGACCTCTGAAATCCCG
AAGGTAGCGCTGTTGAGGAAACAGGGAGCTAAGTCAGTTGAAAAAGTCATTCAAGGTTGGAAAAT
GAACCTGGTTATGTCAGTGGTCTGGCAAGGGCATTGACACAGCTCATATGGCAGCTTCTCAGAATGGCGAAAACCA
TTGCACTGATTGGAACAGGACTGGATGTTTATCCTAAAGCCAATAACGCTTGCAGACTACATGGCAATGACCATCT
GGTCTAAGTGAATATGACCTGGTAAACAACCTCTGAAATTTCATTCTGCCCTAATCGCATCATTGCTGGACTTGT
GTGGTGTGATTGAGCAGAGGCTAAGATGCGTTCAAGGTACTCTATTACGTTGAGCGAGCAATGGAAGAAGGACGCGATGT
CTTGCTATTCTGGTAGGTTAGTATGACTATCAGACGGTTGCCATTGATTCAAGAAGGAGCAAAATTGGTCACCA
GTGGGCAAGATGTTCTGGAAATTGAAATTAA

4181.1 (SEQ_ID_NO. 320)

ATGAAACGTCATTAGCCTGGTCGCTTGTGAGGTTGATGCAAGATTCAACAAACCTGCCCTTCTGGGTATGCAACACTATGA
AACAGTCGAAGCTGTCACCTTGCCTACGGCAACGTCATCACCTCGAAATCAAAATTACTAGAGAAATCGCTAAGAACAG
GGCATTCTGTCACCATATCTCGATATGTCCTGCTGGGACAAATCACTGCTCAGCCAGACTTGTGACGATTCATATTCTCA
CATTCTGACAAGCTCTGTCAGTCAGTCACCTCTCAAACATATCTATTAGTACCTACCGGAAACCAACGGAGATTCCACGAAA
ACTGTATCACACCATCGGAAAGAGTGGTCAACTTGTGACTACCTGCTAGACCTCGCTATTAGAAGTCTGGGAAAATTCACTCCCG
CGTGGCATTCAATCGACCCCTACTACAAACTACGGTAAGCAAGGAAACTAAGTATGAGGGCTGGCAGAACACGCCTTTC
CAACACGACCTTATCCAGAGAAAATTGACAAACCGCTAA

4181.2 (SEQ_ID_NO. 321)

ATGACCGAAACCGTAGAGAAGATAAAAGTAAGTCATTCAATTACTGGGCTGATATCCTCAAGGGGATAGTTCTGGGAGCTG
TCATAAGTGGAAACCGTTCAGTCACCTTGCCTACGGCAACGTCATCACCTCGAAATCAAAATTACTAGAGAAATCGCTAAGAACAG
ATGCTTGGCAACAAATGATACAGTAGTTCTAGGTACGATATCTACAAGTAATTGAGCGAGTCAACTAGTTGTCAGCTCA
GAGTCGGCAAGTACATCTGCATCTGAGTCAGCTCAACCAGCGCTTGCACCTCAGAAGTACAAGTGCATCAGAAATCAGCAA
GTACATCGGCTCGACAAGTATTCTGCATCTACTGTGTTAGGTTACAAACAGCTGCCGCTACAGAAGCAACTGCTAA
GAAGTGTGAAAGAGATGCTAAGAAACCGCTAGTGTAGGTTACAGTACAGTCACTGCAATCTTCAACTCTTATGCTAAG
CGACCGCAAGCGTTGATGGGATTCCATGAGCAATTGCTGGCTTCTATAAAAATGCTGCTGTTTCTGGCAATACGATTGT
AAATGGCGCCCTGCAATTATGCAAGTCTAAACATTGCTAAAGTGAGGACAAAGTTTACAGGTGAAGGTGTAGATTG
GTATATCGTGTCCAATTACTATAATTGAAAGTGACAAATGATGGTCAAAATTGACCTTACCTATACGGTTACGTATGT
GAATCCTAAAACAAATGATCTGGTATATATCAAGTATGCGTCTGGATATTCTATCTATAATTCAAGGTACTTCACACACAAA

4183.1 (SEQ. ID. NO. 322)

ATGGGGCTGAACTTGGTTTATTCTAGCATTGCTGGCTGCCATGGGCTGGTTCCGTTGGAAAGTTCCCTACATGAC
TGCTGCTAATGGCGGTGGAGGCCTTTACTAATCTTCTCATTTCACTATTTAATCGGTTCCCTCTCCCTGCTGGCTGAGTT
GCCCTTGGCGTAGTGTCTGGCTTCGGCTATCAAACCTTGGAAAAGTGGCAAGAAATAACAAGTACAACCTTATCGGTT
GGATTTGGCGCTTGGCTTATCTCTTACAGTGTATCGGAGGATGGATCTAGTCTATCTAGGTATTGAGT
TTGGGAATTTGCTTCAACTTGGTGAACGGGTGATTATGCTCAGTTATTTACTTCAATCATTCAGGATTCGGCTTA
GGAGCTCAAGCGGCCTTATCCTATTGAATATCTCATTTGATCACGTGGGGTCAAAAGGGATTGAAAGAGCTTCGAAAG
TCATGATGCCCTGCTTTATCGTCTTGTGTTTATCATCGGTCGCTCTCAGTTGCCAATGCCATGGAAAGGGGTTCTT
ACTCTCTCAAACCAGACTTTCAAAACTGAGACTGACTGGCTCCCTCATGCTCTGGGACAATCTTCTTGGCCCTCTCAGT
GGGTACAGTCATGGACCTATGCTCTTACTTAGACAAGAAAACCAATCTAGCTCAGTCAGGAACTCCATCGTAGCCATG
AATATCTCGATATCCATCATGGCAGGTCTAGGCCATTCTCAAGCTGATCCCCCTICAATATCCAGTCTGAAGGGGGACCCAG
CTGCTCTTATCGCTTGTGCTCAACTCTTGACAAGATGCCCTTGGAACCATTTCTACGCTCTTCCCTGCTCTTCT
TTGCGACAGTCACTTTCTGCTGTGATGCTGGAAATCAATGTAGACAATATCACCACCAACAGGATAACAGCAAACGTGCCA
AATGGAGCTGTTATTTAGGAATTGGACCTTGTGTTGGCATTCCCTCAGCCCTATCTTACGGTGTCTGGGGATGTTCA
TTTGGTAAGACCTCTTGGCAGCTATGGACTTCTGTTCCAATCTCCATGCCATTGGAGGCTCTACCTTCACTTT
TACAGGCTATATCTTAAAAGGCTTGTGAAATGGAGGAACCTCATCGATGAAAGAGCATGGAAAACAAGGACTGTTCAA
GTCTGGCTCTTCTCTCGTTCTCGTCAATTCCAATCATCATTGTGGCTTCATTGCCAATTATGTAATCAAA
AAGGACTTGAGTAG

4183.5 (SEQ. ID. NO. 323)

ATGTTAAAAAATGGCAGTAAAAGATGTTATCTGCTTGTCTATCTTGTGGGGGTTTCGTTGGTCAGGA
TATGTTGATAATATTCTCACTCTAACACCTCTGGTTGCAGGCCCTTGCCAATGAAATCCTCTCGGTCTGGTGT
ATGGCTGCGCCATTGCTGCCATCTTGTCCAGAGTCAGGAGACTGCAACGATGGAGAAGTGCCTAGCTGCGCTTGTGAAG
TCCTTATGGTAGCCAATTGGTCTAGGAGCTTGTGCTTGTCAAGGTTGGAAAGTGAATTGGTTTATCGTAA
CTAAGAACTGCTATGAAAGTTGGCTCTCTCAACTGCTAATAGTATTGGGATTACGCTTGTAGCTTGTCTATGAATACATT
AAGTTAGGTTACTACGCCCTTCCCTTCCGTTGCTTCCCTGCTTGTAGCTTGTAACTGTTATTCCTGACCAT
CTTGGTTCTGCCATTGCAAACCTATCATCAGTTGCAACTGGAGGAAAAGCATAG

4183.6 (SEQ. ID. NO. 324)

ATGGTCAAAGTAGCAACCCAGACACCGATTATCAGTCCTCTTGCTGTTTATCCTTGGAAACATCTTCATTCCCTTCGATT
GCTCTGACTCTTCGGTAGTCGCATTGTATTCTCTTATGCTCTTACCGTCGATTAAATGTTAGCTTGGATGATCATA
CTTGCCTTACCATCTTTCGAACACTACTGGCAGTTACGGAGATGCTTACAGGCAGTCATGCTTGGAACGAG
GGCTTGTGAGCTTGTATGCCCTGTCTTCTCTGTTCACTAAAGAGCTTCTCTGTACTTGGCTAAAAGGG
GCTATCACGCTTGGCTTATGCCCTGTTGATTGCTTCAATTCTTCTCTCATTCAGCAAGAAATCAAGTCCCTCAAAGA
AGCTTGCCTTATTACGGTCAAGAACATCATTGGCTGCCCTGATTACAGTAAGGTTCTGATGACAGTCTTAAAGTGGC
GCCATCTTACCTGAGAGCTATCTGCTCACGGATATGACGAACATGACAGTGAAGAATAGTATCGGACTTTTATATT
CCTAAAAAAACAAAATTAACTACCTGCTTCTTATTGCTTCAAACCGACTTATTTTATAA

4183.7 (SEQ. ID. NO. 325)

ATGAGAAAGCACCAATTACAAGTTACAAATTAAACCATTATCTATGATGATTGCCCTGATGTAGTCCTAACACCTATCTT
TCGAATTGAGGGATGGCACCGATGTCAGTGACTCAATATTCTAGCAGGAATCATGATGGACCTGTTATGCCTGGCT
ATGGCTACAGTCACAGCCTTATCCGTATGACCAAGGGATTCCCGCTTATGGCTCACAGGAGCGACTTTGGACCCCT
TCTAGCAGGCTCTTTATAAGTACGGTCGAAAATTTCAGTATTCTGCTCTAGGAGAGATTGGAAACAGGTATTGGTT
CCATTGTTCTCATCTGTTATGGTACTCTTACAGGATCAGCTGCTAACAGCTTAGCTGGTTATCACACGCCCTGATTTCG
GAGCAACCTTGTATTGGTACAGCGATTCTTATTGCCCTTCGATTTAATCAAGCAGGAATTCTTAAAGTGCAGGGA
TATTCTTATGTGAAAGGATAGACTGA

4183.8 (SEQ. ID. NO. 326)

ATGAGGAATTACAAATCCCCTTCTATAGGCTCTAGTCCCTCATTCACTGCATTACCAATGAGATTCTGTGAGATGCTG
GCAAATGGGATTGGCTCTGGATGCAAACCTGTATGGCAGATGATCCCGTGAAGTCTGATGTTACTAACGAAAGTC
GGCTCTTCATCAATTGGGCATTGTCAGTGAGAAGAAAAGCAATCCGATGGCAGCTCGTATGCAAACCAATCT
TCTCTCCGATGGTAGATGCGGGTGGCGTAACGACTTCATCCATTGTAAGAGCTTAGTTAAAGACCTTTAGACTATAG
ACCTACGGCTCTAAAGGAAACATGTCAGAAATTGAGCTTGGGATTAAGCACCACGGCCTGGGTCATGCGAGT
GCTAAAGATCAAGAACCGAGGATTGCTTAAGTCTGAAAGACTGGTGCAGACCTATCCTGGTATGCTTTAGTCAC
AGGCTCCAAGGACCTCGTCTGGAAAATCAGGTCGTTACTGGGAATGGCTGACTGAAATTAGACTGGATAACAGGG
ACAGGAGACTTGGTGGAGCTTAACAGCTGTTCTAGCCAAGGAAAGACTGGTTGAGCTCTGCTAGCAGTC
TTATCTAATATCGCTGCTGAGAAAATAGTGTCAAGGAATGGGATTGGAGAATTCTGTTACCAAGTACTCAATCAGCTT
CGCTCTAAGAAGAGATGAAAATTGGTAGATACCATCAAAGGAGAGGTTATGAATAG

4183.9 (SEQ. ID. NO. 327)

ATGAAACATAAAATCGCAATTATCAGATGTTCATGGCAATGCGACGGCGCTAGAACGAGTGTGAGATGCTAAAAATC
AAGGGGCCAGTGAATATTGGCTCTGGAGATATTCTCTGGTCCAGGCAGAACATGACTTAGTCGCTCTGCTAAAGGAC
CTTCCTATCACAGCTGTTGAGGCAATTGGGATGATCGTGTGCTTAGGCTTAGATGGCAATATGCTTAGAAGACC
CACAGGAAGTTCAGCTTGCCTGATGACAGACTATTGATGGAGGCAATGGATCTGCAACGATTGCTGCTACGAAGCTT
GCCTTGCTGGAAAAGAAAATTGACGGATTGCGCTTCTATCTCTATCAATTACCTGACAAAAAAACTATGGTGGTACT
TGCTAGTTGAGAATGATACAGAGAAATTGACCAACTGCTAGATGCGAACGGACGTGGCAGTTATGGTCTGTTCAA
GCAGTTGCTTCGTTATGGAAGTCAAGGGCAACAAATCATCAATCCAGGTCGATTGGCATGCCCTATTAAATTGGGAGGCG
TTAAAAAAATCACGTTCCAGTATGCCGTGATAGAAGTTGAAGATGGGAATTACTCAATATCAATTCTGAAAGTTGCTTAA
TGATTACGAAGCTGAGTTAGAATTGCCAAGTCAAGGGCTTCCCTTATGAAATGTATGAAGAAGTGCCTGCTGACGAT
AACTATCAGGGCACAATCTGGAATTAGCCAGCTTAATAGAAAAGCATGGGTATGTAGAGGATGTGAAGAATTTTTG
ATTTTTGTA

4186.1 (SEQ. ID. NO. 328)

ATGAATGTAATCAGATTGACGGATTATCCTACTTAAAGCTAATAATAGAAAATTAAATGAAACATTATATTGAAAC
CTTGGAAATGAGGCCCTGTTAGAAGAATCGGCCCTCTGCACTAGGTTGACCAACGGGCTTGAAAAGCTGGTTTAGA
GAAGCTCCCAGTATGCGTACTCGTAAGGTAGAGGAAAGAAAATTAGCTAGATTGATGCTCAAGGTGAAAATCCCTA
GAAATTGAAGGAATCTTATCTAAACAGATTGATTATCGATTATATAAAGGTCAAATGGCTACGCTTTGAAATTTC
ACCGAAGATGATTGATTGATTGATTATGCGGAAGATGACATAGCAAGTCTAGTAGAAGTAGGAGAAAAGCCTGAATTCAA
ACAGATTGCGATCAATTCTTAAAGTAAATTGAGATTCTATGAAATTACATCTCCAACTGATATCGAAAGTTCTGGA
ATCATCTGAAATTGGGCATCCCTGATTITATCCAGCTCAGGGCAGGATTGACTGTTGACAAACCGTACCTGGGACT
TATCTATGCTCAAGTCTGGTCAATGAATTAGACATAGCAAGTCTGCCAGAAGTTGACTACTGAATATTCTCCTA
AGTCTGAAAATTCTCTGGTAAAGATAGAATAATGTTGAATTGTTGAGAAGTATGA

4186.2 (SEQ. ID. NO. 329)

ATGAACTGGACCAAGATATTAAAAAAATAGAAGAACAAATCGAGGGCAGGGATTATCCGGAGCCTCTTGGCTATT
AGGACAATCAATGGACAGAGTTCTATTAGGCCAGAGTGAACCCAGAGCATGGCTTGAGACTGAGGAGACTAGTTATGA
CCTAGCTAGTGTAGCAAGGTTGTTGGGTTGGCACAGTTGACCTCTTGTGGAAATAGGTCAAATTAGATATTGATAGAC
TGTTAATAGATTTCACCTGAGAGTGATTATCCAGACATCACTATTGCCAGCTTGTACTCATGCAACAGACCTTGATCCT
TTTATTCTAATCGTATCTTAAACAGCCCTGAATTAAAGGAAGCGATGTTCATCTCAACAGACGAAGTCAGCCAGCCTT
CTTATTCTGGATGTCCTTGGGCTTATTGGAAAAGAATTCTAATCAAGATTGGATGTGATTAAAGGA
TCAAGTCTGAAACCTGGGAATGACGGAAACTTGGGCAAGTCAAGTTGGGAGCTGAGCTGCTGTTCCAAACAGTTAGAGGTTGAG
GCAGGCTAGTGTGATGCCAAGGCTCGTCTCTGGTAGACATGCTGGAGTGTGGTTATTTCGACTATAAAGGATT
ACAAATCTTTAGAACACTATTAGCAGATGATTGCAAGAGACTTAAATCAAATTCTCCTTGGATGACAAGGAAC
GTTCTTAGCATGAAATTGAGAGATTGGTAGACCATACGGGCTATACAGGTACCTTATCATGTGAATCGTCAGAA
GCAAGAACCAACTATTCTTATGAAATCGTACCTATGAAAAGACGAGAGACTCAATGGATATTAGACCGCAATCAAGTG
ATGAACTTGTGATTGCAAGAACAGTAA

4187.2 (SEQ. ID. NO. 330)

ATGATGAGAAGACTTATAATCATATTGGCTGGGAGTCATTCTATAGCATTGCTGTTTCTGTTACTCCT
CAAGAACAACTACCGTGGGAGTGGAACTCCAGGTATTGACATCTGGACGCCCTGGTTTCTGACTCCTTCAATT
TCTCTGGAAACTGGGCAAGTGAAGTACATTGACAATTGTTGGATTCTTACAAAATATCTCAATGCTCTTGTGTTT
TCTCTGATTCTTCAACTCTTATCTATTGCAATTGGAAAACAAAAAGGTCTCTTGTGAGATTGATGTTGTGAGTCTT
GGAAATCGAGTGTACGCAATTATCTGGACTTTCTTGTGATTCTGCAATCGCTCTTGTGAGATTGATGTTGTGAGTCTT
TTGGGTGGCTATCTGGCTGGCTCTTATAAACGATTACATAAAACAGGTAAGGAATTAA

4188.1 (SEQ. ID. NO. 331)

ATGAAGATTCTCTCTTAACTTTGCAAGGCATAAAATTGTTATGTCTGCTTACTTGTCTTCTGCTTGGTTATCGTG
ATGTTTGATGACTTATTCTTCTTGTGATATTGACATTGCGCCGATCTAGCTAAATCGATGGACAAGCAATTAAAATGACTT
TAAAATCAGCATTAGATTTCGTTCTCCAGTCAATCTAGGTTTATCAATCATTTATTCTCAATCATCTGTT

AGGTTTCAATATATTGAGCTAAAAAATAAAGTTTACGATTGAGTATTGGAAGAGAAGTGAGTTATCAAGGGTTAAAAGA
AAGTTGACTTGCAGTTGCAAGTGCAGTATCCCTGTTGATATATTAGTGAUTGTGCTGATAATTGCAATTATAACCTATTCTT
GGGACTTTCTCCTCTGGATGAAATTCTCTATTCTGATGGAAGTGGTTACAAAGACTCCTAGATGGAGAGATAAAAAG
CTATTGTTCTTACTTGTGCTACTAACGGTATTTCATCAATGCAATCTATTGTTACAAATAGTTGATTATGTTGGGAAT
GTGACTCGTCCGGCAATCACCTATTGATGTTCTTGGCTGGTTCTATGCTGTTATAGTGCCTTGCCCTACTATATGGTC
CTATGACGAGTTGATGCAAGCTAGTATGGGATGTAAGTTGATGAAACTCTTACTCCTTATCCTTATATTGTC
ACATGGTGTGAAATATGAAGATAATGTTAA

4188.2 (SEQ. ID. NO. 332)

ATGAAGATAATGTTAAGAATTTTAACATATTGCTAAATAGAAAGATTGTTACTACTTCGTATAGTTCTGATGATGAT
TTTGATAAACCCTATTGCAACAGCGTTCAAAAGCAGGATGCTGTTATCTTCAAGAGAGAATTGATTCAATTTC
CTATAATGACTATTGAGCGAATTAGAAATCCCCAAACTATTGTTAACCTTCGCTTTCATGGTAGGATGGCTCTG
CATTTACTGAAAGTGAATTGGCAGACCATTACCATCACTTGTGCTATCAATCAAGCTCCTTTCGATTATAACAAGGA
AACGATTGGTGTCAATTCTAAATTTTACTCAAGATTGTTGCTGGTTACTTCCTCTAGGAATTCAATTCAA
AACAGTCGCACTTCTTACTTGCTCAGTTATGATGTTGACTTACTGTCTTATCTGATAGCACTGATTGCGGG
CGCTGGTTTCTCTTCTATTGATTGATGGGACAAGAATGGATGATGGATCATATTGTAACAGTGTATTAGT
ACTCTTAAGTTATTAGTTATGTTGATTGTTAGTCGCTTGGAAAGAGAAATTAAAGAAAGGATAA

4188.5 (SEQ. ID. NO. 333)

ATGGGCAAAGGAGAGATGGGCAAAGGAGTTATTGCTTGGAGTTGACTCAGAAGTATTGGTCAACAAGGCTCCAACCC
AATTGGCAAATGGAAAACAGCGACTTCTAACCCAGTATGATAGCAAGACCTTGTGTTGCTGAGTAGATAAGGAAGATAT
CGGACAGGAATTATTGGTATAGCTAAAGGAAGCAGTCAAAGTATGCAATCTTCTGTAAATCTAGCAGGTGCCAGAGT
CCTGGCGAGGATAATGGTAGCAAAAGCAGCGGTGATGAGTTCCAGAATTACAGGGGGAGTTATGGTACAGAGCCAGCT
GTTCATGAAATCGCAGAGTATAAGGGATGTTGACTCTTACTACAAAAAAAGATTACTTACAAAGCTCT
TGCTCAGCAGGCACTTCTGAAACAGGAAACAAGGAGAGTGACCTCTAGCTTACTAGGACTAACAGCTTCTCCTGGT
CTGTTACGCTAGGGAAAAGAGAGACAATAA

4188.10 (SEQ. ID. NO. 334)

ATGTTAAAGTTACAAAAAGTTGAAAAGCTTTATGTTACCTATAGCTATACTTCCTGCAGCAGGTCTACTTTGGGAT
TGGTGGTGCACCTTCAACCCAAACCACGATAGCAACTTATCCAATACTAGACAATAGTATTTCATCAATTTCAGTAA
TGAGCTCTGCAGGAGAGTTGTTACTTGTCAACTTCTCTGTGTTGGGATTATGATTGGCTAGCAGAACAGAGAT
AAAGGAACCGCTGCGTAGCAGGAGTAACGGTTACTTGTCAACTTGTGAGTCAACAGCTTGGTAAACCTTTATGG
CAGAAGGATCTGCAATTGATAGCTGAGTTATTGGAGCATTAGTGTGCGGAAATAGTGGCTGATATTGCAACACCAGATA
CAATATTCAATTACCTCCGTTAGGATTCTTGGAGGTTACGGCTTCAGCTTCTGTTACATCGTCTCTATCTGATT
GGCTTGTCTCTTGTATTTGCCACCTTCAACAATTCTTGTCTACAGGTTGATATATTCTCAGGGGGTCCAATT
GGAACTTTCTATATGGATTTTAATGAGACTTCTGGAGCAGTAGGCTTACATCATATAATTACCTATGTTGGTAA
GAACCTGGTGGTGAACGGTGTGAGGACAAACAGTGGTGGAGCTCAAAAGATTTTGCTCAATTAGCCGATTGG
CCATTCTGGATTATTACAGAAGGACAAGGTTTGCAGGTGCTTCTCAACAATGATGTTGGTTACCGGCTGCCGTT
AGCGATGACCATAGTGTCTCTAAACATGCTGTTAAACATGCGGGTTGTTGGAGTTGCTTAAACATCTTATTAC
CGGATTACAGAACCAATTGAATTATGTTCTATTGCTCAGTCCGTTCTATATGTTGCTCAGGATTCTGTTGTTAG
CTCTTATTGCAAGCCTTAAATATTCAATAGGAAACACATTTCAGGAGGTGTAATGATTCACTTATTGGAA
GCAGGGAAACGCTAAGACGAAATTGGGTTCTCAGATTCCATTGGACTTATTGGAGTGTGTTGATTATATTAGATG
GTTCAACTCTCAATTCAACCTCTAACGCCAGGGCAGGGAGAAGAAGTAGATTCTAAAGAAATTCTGAATCCGAGATTCA
ACTCTAAACTGCGAGATTATTAAACAGGATAGCCTACAAATTATCAGGCTGGGAGTCAAATAATAGAAGATG
TAGATGTTGTGACACGTTACGTTAGCTGTTAGCTGTTAGGAGTTAATCAAGTGTATAAAGCAACTTAAACAAATTGGTGC
AGTTGATGTTAGAGTGAAGGGTGGCATTCAAGCAATCTATGGAGCAAAGCAATCTTATATAAAAGTATTAAATGAA
ATTAGGTGTAGATGATTAA

4188.11 (SEQ. ID. NO. 335)

ATGAAATTAGAAAATTAGCTGTACAGTACTTGCAGGTTGCTGCGGTTCTGGCTTGTGCTGTTGGCAATTCTGGGAAAG
TAAGATGCTCCAAATCAGGTGGTGACGGTGCCTAACACAGAAATCATTGGTGGGATTCCCAGTATTACCAAGAAAAAA
ACTGGTACGGTGTGAACTTATGAAAAATCAATCATCGAAGCGTTGAAAAGCAACCCAGATATAAAAGTGAATTG
GAAACCACATGACTTCAAGTCAGGTCTGAAAAACATCACACAGGCTCGAGCAGGAAACAGCTCCAGACGTACTCTTGTGAT
CACCAGGACGATCATCCAATACGGTAAAACCGTAAATTGGCTGAGTTGAATGCTTCAACAGATGAAATTGTTAAAGA
TGTCAACATGAAACATCGTACAAGCAAGTAAAGCTGGAGACAAGGCTTATATGTTATCCGATTAGTCTGCCCCATTCTAC
ATGGCAATGAAACAAGAAAATGTTAGAAGATGCTGGAGTAGCAACCTGTTAAAGAAGTTGGACAACGTGATTGAA
AAAGTATTGAAAGCACTTAAAGACAAGGGTACACACCAGGTTCTGTTCAAGTCTGGTCAAGGGGAGACCAAGGAACA
CGTGCCTTATCTCTAACCTTATAGCGGTTCTGTAACAGATGAAAAGTTAGCAAAATACAACGATGATGATCTAAATTG
CAAAGGTCTGAAAAGCAACTAGCTGGATTAAAGACAATTGATCAATAATGGTCCACAAATTGACGGGGGGAGATATC
CAAACCTTGCACAGGCTAACACATTCAACAACTCTGGCACCAGCTCAAATGGTATCCAAGCTAAACTTAAAGA
CAAGTAAAGGTGAGAAGTGTAGAAGTACCATCTGGCTTACAGGTTCTCCAGTCCGTAATTGAAAACCTTATGAAAGACA
ACCTAAAGACGTAGTGTGACTCAACTACTCACCATACTACAACACTATTGATGGATTGCTGAAATGAGAACACTTGGTCCAAT
ATAGCGGCTGGACTCAACTACTCACCATACTACAACACTATTGATGGATTGCTGAAATGAGAACACTTGGTCCAAT
GTGCAATCTGTTGACGAAAACCAGCAGATGCTTGAAGCCTTCACTGAAAGCAACGAAACAATCAA
AAAAGCTATGAAACAATAG

4188.12 (SEQ. ID. NO. 336)

ATGCAATCTACAGAAAAAAACATTAACAGCCTTACTGTTATTCAACAATCATTTGCTCTGTTGACTGTGCTGTCATC
TTCCATTCTACTGGATTGACAGGGGATTCAACACTACAACCTGATACAATTGTTATTCCCTCTCAGTGGTCCCTAAATG
CCAACCATGGAAAACCTTCAACACTCATGGTGCAGAACCTGCTTCAATGGATGTTGAAACTCAGTATTATCTCATGGT
AACCATGTTCTTAGTTGTCACCTCATCTAGCAGGTTATGTTGGCTAAACACGTTCTATGGTCAACGCATTCTATT
TGCTATCTTATCGTGTCTGTTGCGCTTCAAAACAGTTGCTCTGTTACCATGGTACGTATGCTCAACTTCTATGGGATCCA
TGACTCTCTGGGAGTTATCTGCTTGTGTTGAGGATGGCATTGGTGTCTTCTCATGAAACAGTTGAGTCAAAC
TACAGAGTTGCTGAACTGCTAAATGACGAGGTTGAGATTGTCACCTCTGGAGTGAGCTTCCCAGTTGAAAC

CAGGGTTGCAGCCCTTCAATCTTACCTTCATCAATACTTGAATGACTACTCATGCAATTGGAATGTTGACTTCACCGTA
ACAATTGACCACATCTCACITGGGGTTCGACCATGCAGGTGAAATGCAACCAACTATGGTTGATTATGGCAGGAGCTGC
CCTGCTGCTGTTCCAATCGTCACAGTCTCCTAGTCTCCAAAATCCTCACACAGGGTATTACTATGGGAGCGGTCAAAG
GATAA

4191.1 (SEQ. ID. NO. 337)

ATGAAAAAAACTTTTCTTACTGGTGTAGGCCCTGTTGCCCTCTTCACTCTGTTTGCCATTGATTCAAGATAAAACT
CTTATCAAGGGGATTGTATATTGACAGACAATACGGCAGAGTTAGACAGAAAGATAGTTACCACTGTTGAGGAGGACTT
TAAGGGCCAATCGTGGACTGGACGTGCTGTAAGATGCCTAGCGGTTGACATTGACCCCTATCCAAAGATTCAAGGCC
GGAAAAAACGGTGCAGAACTAGCAGATGTGACTAGCGAAGTAACAGAAGAAGCGGATGGTTACTGTGAGAGTCTATAAT
CCAGGTCAAGGAGGGCAGACATGTTGAAGTTCGCTCTGCACTTAAAATTTACTTCTTCTTATGATGATATCGCTGA
ATTAAATGGCAACCTCTGACAGATAGTCAAGGTCTATTGAAAAGTTGAATTCTGTAAGGGAGACAAAGGGGGCTGAA
AAACTCTTTCCATACAGGGAACTTTAGAGAGGGACGATTGAAAAGAGTAACCTGATTATACTATCCGTTAGACA
ATCTTCCGGCTAACCGTGGAGTTGAGTTGATGCCATTGGCCTCGGACCGATTGCTAGCGCTAGGGATCAGGATGAA
AGGAATCGTTAGAAGAGTTAAAGATAGAAGAGACTCGATTGTTAGAGAAAAGATCAGAGTAACAAACTCGTTACTTG
GGTCCTCCCTCGATCCTTCCATCTCTGTTATTGAGTGTCTGCTCTTATTTAGAAGAAAAGACACTCCTCAGTC
AAATATGCCAAAATCATCGTCTCTGACACCAATGGAATTAGAGGCTATGGTTTATCAGAAGCAGTCACTCGACCTC
CTTGGAGGAAGTGAAGTCCCTGGCTAACGGAGCTGGAAAATTCACTTGTAGCGCTAGGAGTAAAGAAGATGGTTGCAAGCTT
TAGACCGTGGGAATGTCCTATCATTCAGAAGGAGATGCAGTTGTTGAGGCTAGTAAAGAAGATGGTTGCAAGCTT
TGAGAAAGACTGCCAACATCTAGCTTTCAAGTAAAAAGAAGAAACCTTTCAATTGTTGCGGATTACAAGGTATCTG
ATAGTCTTATCGTAGAGCAGAAAGTTCTGATGAAAACGGATTCAAGCAAGAGGGCTCAACTAACTCTTGAAGA
GGTATTGAAACAGATGCAAGAAGGGAGTGAAGAACAGGTTCCTCTGGGCTCCAGATTATCGTCTTTAAGCTG
GGGAAAAGGCTTGCAGTGGGTATGGGTGCTTGCATCTCCCTTATGGTTGCTTGCAGTGGCTTATAACCTCTACTAAACA
AGACGTTCATGGCTATCTTACCTCCCTTGCAACTACTGTTTCTAGGGTTAGTTGCTGTTCTATTATGGAAAGCTT
CGACTAGATAATCGTAGGGTTCTAAATGAAGCGGGAGCTGAGGCTACTATCTGGACCACTTGTGAAAATATGTC
GTGAGATTGACGATTGGATCAGGTGAAGTATTGTTGCTGGATCGGACCTTGTGCTATGCGACCTTATTGGC
TATGCGGACAAGGTTAGTCATTGATGAAAGGTTCATGAGATICAAGTGGAAAATCAGATATCAATCTATGAGCTTATGG
CTGGCACAGTACGTTTATCATTCAACAGCACAAATGAGGCAATTATGCTAGTGTGCAAAATACAGCAAGCACCTACTCTGAT
CTTCTGGAAGTGGAAAGTCTGGTGGCTCTGGAGGGGGAGGTGGCGGAGTATCGGTGCCTTAA

4191.2 (SEQ. ID. NO. 338)

ATGAAAAAAAGTAAGAAAGATATTGACAGGGCAGTTGCAAGGACTGCTGTATATCTCAGTTGACAGCTTTCTCGATAG
TTGCTTCTAGCAGAACACGCTGAAACAGTCCAGCGTAGGAAAAGTGTGATTAAGGAGACAGCGAAGGAGGAGCGCTTC
TAGGAGATGCCGTCTTGTAGTTGAAAACAAATCGGATGGCACAACGTTGCGAAAGGAGACAGGGCGAAACAGGGAG
CGATATTCTCAAACATAAAACCTGGACATACACCTTGACAGAACGCCAACCTCAGTTGTTATAACCTCTACTAAACA
ATGGACTGTTGAAGTTGAGAAGAATGGTCGGACACTGTTCAAGGTGAAACAGGTAGAAAATCGAGAAAGAGGCTCTATCTG
CCAGTATCCACAAACAGGGACTTATCCAGATGTTCAAACACCTTATCAGATTATAAGGTAGATGGTCTGGAAAAAACCGGA
CAGCAGAACGGCTGAATCCGATCATGAACTGTTGATCCAGAGTCAACTTCAAAGAGAATTATCAAGTGAATA
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GGATGTCGTTATCTGCTGATACTCAAATAGTATGAGTAACATTGAAACAGGATGCTCGAGTGGCGAAAGAGCTGGT
GAGGCGACACGTTCTTATTGATAAAATTACATCTGATTGAGAAAATAGGGTAGCGCTGTGACTTATGCTTCACTATCTT
TGATGGGACCGAGTTACAGTAGAAAAGGGTAGCAGATAAAAACGGAAAGCATTGAATGATTCTTTTGGAAATTAT
GATCAGACGAGTTACCAACCAATACCAAAAGATTAGTTTAAAGCTGACTAATGATAAGAATGACATTGAGAATTAA
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CCAACATATGTCGATCCGATTAATTTAATCATGCTACGTTGCTCCATATCAAATCAACTAAATGCAATTAGTAA
TCTCCTAATAAGATGGAATACTATTAGTATTACGCAAGCAACTAGTGGAGAACATACAATTGACGGAGATG
GGCAAAGTTACCGAGATTTACAGATAAGACAGTTATGAAAAGGTGCTCTGAGCTTCCCAGTTAACCTGAAAATA
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TGCTCCTGATGGTATGATGTTACCGTAGGTATTGTTAAACGGAGATCTGGTACGGATGAAAGCAACGGCTACTAGTT
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AGAGATGTGACTTCATCGTCCACAGATATACAGCGGGTTACGAGTTACGAAATGATAAGCAACTATTACCAATGAC
CTATTCCCTCAAAGAGAGAATACCTCGAACCTGGTATCGGAATGTTGCCATTCTATGATAGGTTGCATGATGATGGGA
GGAGTTCTATTATACACACGGAAACATCCGTAAC

4191.3 (SEQ. ID. NO. 339)

ATGAAATCAATCAACAAATTAAACAATGCTGCTGCCATTACTGACAGCGAGTAGCCTGTTTCACTGCAACAGTTT
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TTAGAAACAGTAACATGCTGTAATAAAGGGTGTCTACCTGCAATGCAAATGAAAAGAAATTGCGGTCTTATGTTGCTT
GGACAAATACTAATAATGAAAATTGATGAAAATGGCCTAAACTCTAGGAGTGAATATTGATCCTACACACATTTAAACTCTC
AGGGCAATGCCGGCAACTGCAATGAAAATTAACAGAACGCTGAAGGAGCTAATTAAACACGGAAATTACCAAGCTGC
TAAGTATAAAATTGAAATTCAACAGTTATCAACTTATGTCGGTGAAGATGGAGCAACCTTAACAGGTTCTAAAGCAGTTC
CAATTGAAATTGAAATTACCATGAAACGATGTTGGATCGCATGTTGCAACAGGTTCTAAAGCAGTTC

TAAAGATTCAAAGGTAAGCAAATCCAGATACACCACGTGTAGATAAAGATAACACCTGTGAACCACCAAGTTGGAGATGTT
GTAGAGTACGAAATTGTACAAAATTCCAGCACTTGTCAATTATGCAACAGCAAACCTGGAGCGATAGAATGACTGAAGGTT
TGGCATTCAACAAAGGTACAGTGAAGTAACGTGTGATGATGTTGCACTTGAAAGCAGGTGATTATGCTAACAGAAGTAGC
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GATCCAACAGAGCCAAGTGTACATATGGTAAAAGTGTCAAAGTAAATGATAAAAGATAATGTTAGCTGGGAG
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AAAGAGAAAGTGTACAAAGCTCAAGCTGTTATAATGCTGCTGATTGCTGCCAACATGCAATTGAAATGGGTGGCAGATA
AGGACAATGAAAATGTTGAAATTAGTTCTGATGCAAAAGGCTGTTGAAATTACAGGCCCTTGTGAGGTACATATTAC
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GACTGGACAAGGATTGGTAACTGCTGGTCAAGTAAAGATGACGCTACAAAGTGTCAACAAAAAAATCACTATCCC
ACAAACGGGTGGTATTGGTACAATTATCTTGTGTAAGCAGGGGCTGGATTATGGTATTGCAAGTGTACCGCATATGTTAAA
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4191.4 (SEQ. ID. NO. 340)

ATGACAATGCAGAAAATGCAGAAAATGATTAGTCGTATCTTCTTGTATGGCTCTGTGTTTCTCTGTATGGGTGCACA
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GATGGTCACTGGTTGCAAGTATGGAAGTGGATGATTGTTATCCCTATGATGATCGGGTCAAAATTGTAAGAGACTTGCATTC
GTGGGATGAGAATAAACTTCTTCAAAGACTTCGTTGAGATGACCTTCCCTGAGAATCAGATTGAAAGTATCTCATA
TTCAAATGGCTTACTATGTTGCTCTATTATCCAGACGGATGCGGTTCTTATCCAGCTGAATTCTTTGAAATGACAG
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TGGAGAATACCGTTACAGTTCTCTGGTCAAGTAGGGAGAACTCTCTACTGATAAAAATGGAGAGATTGTTGACAAAT
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GGTAGATCATCAGCTGGTACGGTTGCAATCAGAAATTACCAAGTGGCAATGTTGACTTATGAAAGGTGGATGGT
CGGACCAATACCTCTTCAAGGGGCAATGTTCAAAGTCATGAAAGAAGAAAGCGGACACTATACTCCTGTTCTCAAATG
GTAAGGAAGTAGTTGTAACATCAGGGAAAGATGGTCTGGAGTGAAGGTCTAGAGTATGGACATACTATTATGGG
GCTCCAAGCTCAACTGTTATGTTCAATTAAACATGCCCTTCAACATCGGGAAAGATACTCGTAAGGAACGGTAA
CAGTGGTAAAATAACAGCAGGACCAACGGATTGATGTGCCAGATAACAGGGGAAGAAACCCCTGTATATCTGATGTTG
CCATTGTTGGTAG

4191.5 (SEQ. ID. NO. 341)

ATGAGCCACATACATCTATTCACAAGTCTTGTGATGCTAGGACTTGTCAATGTTGCTCAAGCCGATGAATATTTA
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CCAATCAGTATGCCAACGGATACGATGTTCAAATGCCAAGAAAATGCTAAGGACTTAGTAAAGAACCTTGGTTAA
AACCAAGTGGAAAGGTCTAGTCCCTGCCCTACTTCTGGTAAGATTGACATGATTATCGCAGGTATGAGTCCAATCGCAGAA
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GTGCTAAATCTTGGTACCTTAAAGGTGCAAAAATCACTTCTCAACAAAGGGGCTACCTTATAACTGATTGACAAATC
CCAGGTGCTAAAAGAAAACGCCATGGGAGACTTCGCTCAAATGCGACAAGCTTGTGAGGCTGGTGTATTGATGCTTATG
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TGGGAAGAAGATAACAGTATCGCTATCGGCTTGTAAAATGACAATGTTAGCCAAATCAATGCCAGCATGAAACC
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GCACTAGTTCTTAGCCAAGTTGCTAAAATTCTTCTGAAAACCTGCAACAAACTCTTGTGCTGGTGTACTCTTAA
TCTCTATGTCGGAACCATCATAGGTTCTTAAATGGAACCTGGCATTGGTGTCTCGTACTGCTCTCTCTGAAAACAAAG
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TCGATGGTTACTATGGAACCTGCCAAGCTTGGGATCAAC

TTGACCGTACACTGGCTGCTATCTCATCGTTCAATCAATACCGGTGCTACATGACTGAAATCGTCCGGTGGTATCTA
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AGTAGTCCGTAACATCCTACCTGCAACTGGTAATGTTGATCATCAATATCAAAGATACTGTTATTGAAACGTTATCTC
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TTACTTGTCTCACCTCACCGTAACACGTTACGCTTATGAGCCAGAATGGACATGGGACACTACACTACAGGTG
CTAACCAAATGCAAACGGAGGATTGAAATAA

4191.6 (SEQ. ID. NO. 342)

ATGACACAAGCAATCTTGAATTAAACACCTCAAAAATCTATGGACAAAACGAAGTGCTAAAAGACATTCACACTG
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AACACCAACTGATGGCAAATCCTTATCATGGACAAAAGCTCCTCGAAAAGGCTATGACCTCACGCAATACCGTAAAAG
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AAAACCGAACGCACAGAAGCTGAAAAGGAAACCTGAAAAGGCTGGCATGGAGAACGCTACTGCCAG
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CTTGTGAAACCAACATCAGCTCGCAGGAAATGGTTGGAGAACCTCTCAAATCATGCCAGGACTGGCTCAGGAAGGC
TTGACCATGATTGTCGTAACCATGAAATTGGAATTGCCCCGTGATGTCCTCACCGTGTATCTTATGGATAAGGGCGTGT
CGCTGAAGAAGGTAACCCAGAAGACCTCTCACCAATCCTAAAGAAGACCGAACAAAGAGTCCTTCAACGCTATCTCAA
ATAA

4192.3 (SEQ. ID. NO. 343)

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GTCATGATTGGTTCTTGTAAAGACAGGCCATGGTATCTTCTCCGATTCCAAGAAAAAAATGGCTTGTATTGATTT
GACAGTATTAGTGTCTAGTGTCCAGATCTTTAACGTTCAGACAGCTAAACATGTTAGTCAGTCAACTGCGGAAGGTTGGCTG

TATTGATTGGTTAGTGGGACTAACCTTGAGAGCTAGGTATTTATAGCCCTGTTCTGGTTCACGTGATGGAAGAA
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CTCTCCCTCATTTCAAGCCTGCCTAGTGTAGATATCTCGTCTTGCAACAGTGGAAATCATCTTGCTGGTTGACCC
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4193.1 (SEQ. ID. NO. 344)

ATGAACAAGAAAACAATGGCTAGGTCTGGCTAGTTGAGCAGTGGACTTGCTGCATGTGTAACCGCTCTCTC
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CCAATCAGCTGGAAAGGTTGAGGCTTGGTAAAGAACACAATCTTCAAAAGATAACGGTTCACTTACTTCCAATCA
ACAAGTGAAGCTGACTACGCTAACAAACTTGCAACAAAGCGCTGGAAACGTTACAACCTAATCTCGGTGTTGCCCCITA
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TGTGCGAGCGTAACITCGCTGATAATGACTCAGGTTACCTTGAGGCTGAGCAGGATTAAGGCTGGTGTGCGTCACTGAGACCC
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GCCGCGTGCAGATATTGTTACCAAGTAGCTGGTACAGCTGGTACAGGCTGGTACAGGCGAAATCTCTAACGAAAGCC
GTCTGAAAAGAAAAAGTTGGTATCGGTGTTGATCGTGAACAGGAGAAGGAAATACACTTCTAAAGATGGCAA
AGAATCAAAACTTGTCTGTATCTACTTGAACAGTGGTACACTGAAAGGATATTCTAACAGGAGAAGGAGA
GAATTCCCTGGCGGTCAAGTGATCGTTACTCATGAAAGGATAAGGGGTTGACTGGCAGTAACAAACCTTCAGAAGAAG
GTAAAAAAAGCTGTCGAAGATGCAAAAGCTAAATCCTGATGGAAGCGTAAAGT
TCCTGAAAATAA

4193.3 (SEQ. ID. NO. 345)

ATGTCTAAAAAATTACAACAAATTICGGTCCCCITGATTCTGTATTCTAGGAATTITACTCGGAGCCATTGTCATGTGGATC
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TGCTATGGGTCTCTGGTCTGATTGGTCTTGGCTTGCCTGAGCTGAGCTGGTTCTTAAACGTCGACTTCTGGTCA
GGCTTGGCAGGTTGGATTCTAGTGGTGGATTGCGCATCCAGATATGCCCTGGCTTGTGATGATTCTAGCAACCA
TCGTGATTGCCCTGATTGCTGGGATTGCGGAGCATTCCAGGTTCTAGGGCCTATCTAGGGACCTCAGAGGTTATT
GTAACCATCATGATGAACTACATTGCTTGTATGTTAGGGAATGCCATTATCCATGCTTCCCTAAAGACTCATGCAAGTAC
AGATTCGACCATTCGTTGGGCTAATGCAACCTATCAGACACCTTGGTGGCTGAGTTGACTGGTAACTCAGGATGAAT
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GTGGTCTTAAATCCACATGCTCAGAATATGCTGTTATTCTGCCAAGGGACTATTATCCTATCTGATTATTCTAGGTGCC
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TAACGGAATGGCGGTAGTTGCTTGGCGCAACTCACCATTGGTATACTCTTGCAAGCTTCTTATTTGGCTTCTCCAAGT
TGGGGCTCTGGTATGAATGCCCGCAGGTAACATCTGACCTGTCAGCATTGTAACAGCTTCTTATCTTGTCACTG
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4194.1 (SEQ. ID. NO. 346)

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TTGATATTAGTATCGGAGTGGGATTATCTCTTACGGCTTGTGATTGTCAGCTCTTGCAGCTTCAAGGTGCAAATGG
TGGCTTCTAGGAAAATGAGGAGCTCAGGCCACGCTTAAGGCGCTCGAGAACAAATATCCAGGTGAGGATATGGAAGCA
GAACCAAACATAGAGCAGGAAATGCTTAAAGTATTAAAGGAAATGGGTGTCAGACAGTCAGACTCTTGGCGATTGTT
TCAGATGCCGTTATTGGCCCTGTCAGGCTATCAAGAGGTTGACTTTAAAGACAGGTCATTCTTATGGGATTAAACCT
TGGTAGTGTGAGTACACCCCTGTTCTCCGATTAGCAGCAGTATTACCTTTAAAGTACTTGGTTGTCACAAAGCTT
GTCTGAGCGAAATGGCGTACGACTGCGATGATGTTGAGGATTCCAGTCTGATTGTTATCTTGCAGTTATGCCAGGTG
GAGTCGCCCTACTGGACAGTGTCAATGTTATCAAGTCTGAAACCTATTCTGAAATAATCCATTCAAGGATTATCGCA
GAGCGCGAGGCCGTAGTACAGGACAAAAAGATTGGAAATAGAAAAGCAAGAAAAGGCTCAGAAACGAA
ATAA

4194.4 (SEQ. ID. NO. 347)

ATGGTTATCGATCCATTGCTATCAACGAACTAGACTATTACTAGTTCACACTTCCACAGTGTATCATATCGACCCATACAC
AGCTGCAGCAATTCTCAATAATCCTAAGTTAGAGCATGTTAAGTTATCGCTCTTACCACTGTGGACGAATCTGGAGGAT
GGGGTGTCTTCAAAAGAACGTTATCGTGTAAACCAAGGGACTATCGAATTAAAAGATATGAAAGATTCATGCACTGAG
ATCATTTGACCGTACTTGGTAACTCTCCAGTGAACCGGTGCTGATGAGAACAGCGGTGAACTTGTGGCTTGGCTGTTA
CAGATGAAGAAAATGGCTTAAAGGCTTAAACTATCTTGAACACCCAGGTGGAACCATCTATGGTCACTGAGTCTCA
CTTCTCAAACATTGGCAAACATGGTAAAGACTTAAATTGATGTTGCTTGAATAACTATGGTAAAATCCGTAGGTA
TCCAAGACAAAATGACATCTATCGACCTTCTCGTATGGCAGAAAATCTCGTACCAAGGCTTATCCAGTTCACTATGAT
ATCTGGTCTAACTTCTATGGCTTCTACTAATGAGGATTCTAGAACTTGGAAAATGCCAAAAGATCGCTTGCACATCGATTCCA
TCCATTATCTGGGAACTGGCGGTAAAGTACACTTCTCAAGATCAACACTTACTAGTAAAGTACATTCA
ATGATTGTTGAACAAGACTCTAACATTAAAGCTTGTCTATAA

4196.2 (SEQ. ID. NO. 348)

ATGTTCCCTTCAGGCTGGTTGTCTAGTTGCTAATACCTATATCCATGATTTACTGGGGTTCTTCCCAGATAGTCCATT
TAAATGCCCTTGAAGTGTCTATTGGCTCTTGGTAAAGAACCTTGAAATTATTGTCATTGTTTGTGTTGGCTTGA
TTCTGTGCGAAAATTAAATCTTGTGTTTACTTGGAAATTGCTTCCGGTTGGGATTCCAATGATTAAGGATATTGGTTATA
TTCGTACGGATTGCGCAGGGCTTGTACTTACTATTCGCAATTAGAGCGTATCATCTCAGGAATTGCGCTCTCACTGG
ACTTTTCTAGGTCTAGTGTAGTGGTTACTGCTTACAGAGGCTATAAGGAGCAGAAGGTTGGCAAGAAAAGGCC
TTATTGTTCTAGGTTAGCCTGGGAACTCACTCTGTTAACTCTCCTTGTGAGGTTGGAAACAGAGTGTGCTTACGGA
TTCCAGTGGTACGGCTATTGCTCTATGGTTTATCATGCTTATTGCTTGTGAGGAAACACAATGAGTTGATGACCTAG

4197.1 (SEQ. ID. NO. 349)

ATGAAGGTGAAACCACCGTGCACGTCCTTCGAGGATGTCGCACTTTTATTAGGATACTAATTATGGAGTTGCAAGAATT
AGTGGAGCGCAGTTGGCAATCCGACAAGCTTATCACGAACGACTGGAAGCTAACCATCATGATTCAGCTGACGGTAGAAGA
AGACCTCTGGTTATCTAATGATATTGAAATTCCAACGACTGGTATGACAAAGCAAGGACGCTACTATGATGAAACA
CCCTACACACTGGAACAAAACCTTCAGAAAATATCTGGTGGCTATTAGAACCTTCTCACCGTTGGATATAGACATTCTGAC
GGAAATGGAAAACCTCTCTGATAAGAAAAGCAATTGAACGTTAGGACTTGGAGTAG

4197.4 (SEQ. ID. NO. 350)

ATGCTTGATTGAAACAATTCTAGCCTATCTGCGCTCCGAGTCGCTTTTATCTATCTGCTTGGCATTTCTTG
TCTTAECTCTTCAGTTTATTGCCAGTCTAGGAATTACTCCCTACTTTCTCTTGTTGTTGTTGAAACCATAATT
TTCACTTGGGACATATTGGTGGAAACCGCAGGCTATCGCAGGAACTCTCTATGAGGAGAGGGAAGCCAAGTCCCTTGG
AAATAGCTTCTAGCAGAAAATAGAAGCGCTGAGATGGAACCTATCAGCAGAGGTCAAAGCAGAAAAGAAAAGTACGG
ATTGCTGTTACTATACCTTGTGGTCCATCAGATAAGACCCCCATTGCAAGCAGTCAACTTGTGAGAAGTGGT
GACCGCCAACCTGAAGCAGCAGCTAGAACAGGAATTCTAAACTGACTCCTATACCAACCTAGTTTACAGTACCTGCGTT
TAGAAAGTTCCATGATATTGGTCTTAAAGCAGGTTCAATTGAGGACTTGGTCAAGGAAATAATTGTAATATGCTCTT
TTCTTATTCAAAAAGGTTAAATGCAATCTACATGACCTTGTAAAGAAATCGTACGGATAAAAGTGGCTGCTAGTGG
TTATTGAGGAAATCATCTAAACAGTCTCAAGTACACCAAGGAAGGGTGTGGAGGATTTATGGATGACCAAGAGCTTGG
TATCAAAGATCGGGATCGGGAAAAACAGTGTGCTCTCGAGATTGAGCTGGCTTTCAGGATACAATGGCGT
TTGACCCAGCAGTCTCTGGACTTGGCTTATCTATCTAGAAGAAAATTCTGAGAAGAACGGGGACCAGATCGTATCGAGTC
TGAGTCGGAAAAGGAACGACAGTGGGATTCAAGTGTCAAGTGAACCTAGTCTGAGTAA

4211.2 (SEQ. ID. NO. 351)

ATGGAACCTTAAACACACAATGCTGAAATCTTGCTCACTGCGAGCTAATAAGTCCCACATCCCGAGGATGAACTGCCAGAGA
TTGCCCTAGCAGGGCGTTCAAAATGTTGTAATCCAGCTTATCAACACTATGTAACCGTAAGAATTCGCCCCATCA
GGAAAACCTGGAAAACCCAGCTCTGAACCTTTAACATTGATGACAAGATGCGCTTGTGATGCTGGTATGGCTA
TGCTCGTCTTCTAAAAGGAACGTGAAAAGTGGGGGTGATGATTGAGGAGTACTTAACGACTCGGAAAATCTCGTGC
GTGTCAGTCTAGTTGACCTTCGTATGACCCGTCAAGCAGATGTGAGATGTAAGAATTCTCAAGTATTATGAGATTCC
AGTCATCATGTGGCAGCAAGGGGACAAGATTCCTCGTGTAAATGGAACAGCATGACATCAGCAATCAAAGAAAATT
AAACTTGACCCAGTGTGACGATTCTACCTCTTTATCTGCACTGAGCAGGGATGGTGGAGGCTGGATGCAATCTTAG
AAAATTGTGA

4211.3 (SEQ. ID. NO. 352)

ATGACAAAGAAACAACCTCACTTGGTGTGACAGGGATGAGTGGCGCAGGGAAAAGTGTAGCCATTCACTGCTTGCAGG
ATCTAGGTTATTCACCAATTGATAATATGCCGCCAGCTCTTGCCTAAGTTTGTGAGCTGGTGAATTAAAGGAAGACAAT
CCTAAGTTGGCTTGGTAGTGGATATGCGTAGCCGTTCTTCAAGGATTGAGCTGTTGGATGAGTTGGAAAATCA
AGATGGTTGGATTTCAAAATCCTTTGGATGCGGCTGATAAGGAATTGGTGTGCTGTTACAAGGAAACCAGACCGAGT
CACCCACTAGCAGCAGACGGTGTATTAGATGGAATCAAGTGGAACGTGAACTCTTGGCACCTTGGAAAATATGAGCC
AAAATGTGGGATACGACTGAACTCACTCCAGTGAGTGCGCAAACCCCTTGCAAGAGCAGTTTCAAGGACAAACAAGC
CCAGITCTTCCGTATCGAAGTCTGTTGGATTAAAGTATGGAATCCCGATTGATGCGGACTTGGTCTTGTGATGTCGTT
CTTGCCTAAATCCCTATTATTACAGAACTGAGGAAACCAACGGGTGTGGATGACCTGTTATGATTATGTCATGAAACCATC
CTGAGTCAGAAGACTTTATCACATTATTGGCTTGTGAGCCATTCTGCAAGTACCAAAAGGAAGGTAAGTCCGTT
TTGACCAATTGCCATGGGATGTACGGGTGGACAACACCGTAGTGTGGCATTGCTAAACGCTTGGCAGGACTTATCCAAGA
ATTGGTCTGTTAATGAAGGGCATCGGACAAAGACCGCAGAAAGGAAACGGTAAACCGTATG

4211.4 (SEQ. ID. NO. 353)

ATGAGAAAACCAAAGATAACGGTGTGAGGGACTGGAAGTCCGTCATTCTAAAAGTCTGGGGAAAAGATGTG
GAAATCGCAGCTATCGTACGGTGGCAGATGATGGTGGTCTTCAGGTGAACTCCGAAAATATGCAACAGTTGACACCGC
CAGGTGATCTCGTAATGCTTGTGGCATGTCGGATATGCTTAAGTTTATGAGAAGGTCTTCACTGTTCTGAG
GATGCGGAGCCTTGTGGCCATCATTGGGAAATCTCATCATTGCTGCTGTCAGAAATGAGGTTCAACCTATAATGC
CATCGATTATGAGCAAATTCTCATACAAACAGGAAAATTCTCCAGTGCACCATCCTTGGACCTTCACTGCGTCT
TCAGGATGGGAGCAGAAGTGGCTGGAGAGACTCATTTGAGACCATCGAGGCTATAATTGACAATGTCTATGTGACCAATGCC
CTAACACGATGATACGCCCTGGCCAGCCGTGAGTAGTGCAGACCATCCTTGGAAAGTGCACATGATTGCTTAGGGCCAGGTT
CCCTCTTACCTCTATTGCCCATACTGTAAGGAAATTGGGGGGCTCTTTGGAAACCAAGGAGAAATTGCTTAT
GTCTGCAATATCATGACCAACGTGGGAGACCGAACACTTACAGATAGCGACCACTGGAAGTCTGATCGTACCTTG
GTGCCCCCTTATCGACACTGCTTGTGTAATTGAAAAGTGGCTCAGGAATACATGAAATTCCAACCGTTTGATGAAATAC
TTAGTCAGTGGAAACACGATTGGTAGGTTAGTGTGAAAGCAAGTTGGCAGGTTGATGCCATTACAGGTGAAAAA
TGGCGGTGCTTCCACGATGGAGATTGATTTGGAGGAGTTGATGCCATTACAGGTGAAAAATGA

4213.1 (SEQ. ID. NO. 354)

ATGAAAATTGATAAAAGGATGTTAATTAATTAGATTGATTGTTAATTAGCAGACAGTGTATTATATAGTAGCATTGGCA
CGTAGCAATAATTATCTCGAGCATGTTCTAGGAATTATTGCACTGAAATTATCTACCGGATTGTTACTAATCTT
TGACCGATTGAGCAGAGTAATCCGAAAAAAATTCTTATAATATCAATTGTTGTTCAATTAGCAGTGGCTGTAATT
TATTATTAAACCAATATCATTTGGGTGATAATGAGCTAGTGTGTTATTGAGCTAATTGCAATTCTCTTTAGTATTGTT
ATAGAAGATGTGTTGATTCTCAAGGGTGAATATGATAAGATTGATTGCAATTCTCTTTAGTATTGTT
ATTAGATTCTTATTTAAATTGAGCTATTGTTAAAGTAAACTAGCAATGCAATTAGAAAATTCTCTTCAATT
CAAGAGAGAAGTGTGCAAGGTACAAAGTTTAAATAAAATTATTATAAAACCGATATTCTTAACGCTTAA
ACTTTTATTCTATTGAGCTAGTGTACCGATTGTTCTATTGAGATTGTTAGGTTAAGTGGTATTGTT
ACTATTGCTGGTTGGGTGATATTGGGAAATATGCTAGCGCAATCGTAATAAAATTGAGTGGTCCGATTGTT
TGTTTTCTTTTGAGCGCTCAAGTGGTTAGTAGCAATTGTTATAAAAGACTATACTTATCACTTATTGTT
TGTTTTATGCTAAAGGAGCTTCATAATTGTTAATTGTTGACCAACAAACCTCCACATCAACTCTTGGTAGGGTA
AATACTACCATGTTCTATTATCTTGGAAATGCCAATTGGTAGTTAGTGTGAGGAGCTTATTGATTGAAATTGAA
TTAGTGTAAATTGCTATTAGCATACCTTATTGTTGTTCTTATATCTGAGGATAATGGATTGAAAGAATTGAT
ATTAG

4213.2 (SEQ. ID. NO. 355)

ATGATGTCTAACAAAATAAGGAAATTCTGATTTGCATTCCTATACAGTCCCTTATGTTGATGGCGTTAATTGCTG
GCTCTTAAATGCCATCTGCCATTGAAATTATCTGTTATGTTAGCTCTATATGGCTCCTTCTGTCAGGATAGAT
TGATCCAACAATGGAAGGAGATTAGAAAGACTAAAAGAAAATTCTCTTGGAGCTTAACAGGATGGCTCTTCTCATTCT
GATGACTGTTGCTTGAATTGATCAGAGATGTTGAAGCAGTTGGGACTAGATGGACAAGGCTAAATCAGTCAATA
TTCAAAGTACCTTCAAGAACCAACCAACTGATAGCTGTTGCTTGTGTCATTGGACCTCTGGTAGAAGAATTATTTC
GTCAGGTCTTATTGCAATTGAGGACGGTGTGAGTTACTAAGCATTATTGAGGACTTGTGCTGACTC
ATATGCACAGTTGGCTATCAGAGTGGATTGGCAGTGGTACTTAGGTGGAGGCCCTGCCCTTCTATTATATGTGA
AAGAAAAGAGAATATCTACTATCCCCTACTTGTACATGTAAGAACAGCCTCCTTAATCATTAGCTACAGTATA
GTAAAATGA

4224.1 (SEQ. ID. NO. 356)

TTGAAAAAGCCAATTATCGAATTCAAAACGCTCTAAAGTTTGAAGACAGCAACACCAAGGTTCTCAAAGACATCAACT
TTGAGTGGAGAAGGAAATTCTACACCCCTAGGTGCATCTGGTCGGGAAATCAACTATCTAAACATTATTGCAAGG
TTACTGGATGGCAGCACAGGAGATATCATGCTAGACGGTGTTCGATCAATGATATTCCAACCAACAAGGGCAGCTACAT
ACCGCTTCAATCCTATGCCCTTCCCACATATGAATGTTGAAAATGTTGCCTTCACTTCGCTGAAAATTGAT
AAGAAAAGAAATCGAGCAGCGTGTAGCGGAAGTCTCAAGATGGTCAGTGGAAAGGTTATGAAAACGTTCCATCCGCAA
CTTCTGGAGGACAACGTCAGCGTGTGGCATCGCCCGTGTATCATCAACCAACCCGTGTGGCTTGGACGAGCTT
ATCAGCGCTGACTTGAATTGAGAACAGACATGCACTGAGTGGCAATTACAACAAACGATTGGGATTACCTTGTG
TTTGTCACTACGATCAGGAGAACGCTTGCCTGAGTGTGAGCTTGTGTTGATGAATGATGGCAGGATTGTCAGCTGG
AACCCCTGTGACATCTAGTGAACCATCAACCAACTTGTGCCACCTTATCGGGAGTCACACATCTGCCAGGACCA
TGATTGAGGACTACTTGTGCAATTAAACGGCAAACGCTTGAAGCGGTTGATGGGGATGAAGCCAATGAACCTGTTGA
GGCGTTATTGTCAGAGGACTTGGCATTACCCCTCTGAAGAAGGCAAGCTCAAGTTAAGGTCGATACCCAGCTTCC
GTGGAGTTCAATTGAATTATGCCCTATGACGAATTGGAATGAATGGATGATCCACTCAACCGTAAGGCTATCGTGG
TGAGGAATCGGTCTGGACTTGAACCAGAACATCCACATCATGCCCTCAATGAAACCGAAGAAGAGTTGATGCTCG
ATTGAGGAGTACGTAGAAATCGAAGAGCAAGAACAGGTTGATCAATGCAATCGAGGAGGAAGAGATGAAGAAAACAA
GCTCTAA

4252.1 (SEQ. ID. NO. 357)

ATGAAATCAATGAGAACCTTATTGTTAGCTTAATTCAAATCAGTTGAGTACCTGTTCTATGGAAGGAATGCATCTT
GTCCTTAAACAAAGTACAGCTTTTCATCGGAAGCATGTTTCAGGAATCTGCTGGAGTAATTATCTTATAC
CCGAAGAAGAAGTCCATAGTGTCTAGCCAGTAAGAAGTGGTGAAGCTTTACAGTATGTTACTCTTAATTAAATTG
TAGGAGCTGTTCTGTTGTCAGATAACTTGTICATCAAAATACGCTGCAGCAAGAATTAGTGTGACTTTTATGCCATCCT
TCTTTCTTCTATTGGCTAGATTGCTGATTTTTACCTTGAAAATACGTGCGCATTGCTATGCTGGACAGAA
AAAAGACAGTGGTGTGACTATTAGCAACACTCTTTCTTAAGAAATCCAATGACCATTGCTCACTCTGATTATATTG
GACTGGGCTGTTCTGTCAGCCTATCTGTCCAAATTGCTTAAGAAGGAAGTTCCTTATGGTCATATTCCGAGATC
TTGTATTGGTCATTGTTACGCTCATTTCTTTAG

4252.2 (SEQ. ID. NO. 358)

ATGGTTAAAAAAATTATTGGAATGGTCTAGCTTACTTCTGTAACTGAGTAGGGAGTAGGTGTTTGCTTACTATT
CAACAAGGGACAGAACCTTCTAGCTAAACCTATAAAAATCGCTGAAGAACCAAGGTTATTGAAGCGACTGAACCTCTA
ACCATTCTGTTAATGGGAGTGGACACCGGAAATGTTGAACGAACTGAAACCTGGTGGTAGAAGTGTGATAGCATGATCTG
TGACAGTGAATCTAAAAGAAAAACAAACAAATGAGTGGTTAGAGCGGGATATTCTGACCGCGATTGAATCAGGGATG
GTCAGGCTCATGAAGCGAAACTGAACATCAGCATATGCAGATGGTGGAGCAGAGCTTGTATAGAAACCAATTCAAAATGA
TGAATATCCATATTGATCGTATGTGTCAGGATTGCAATATGAGAGGATTGCAAAACTAGTGGATGCACTGAGTAGGAGGTATTACAGT
CAATAATATCCTAGGTTCCAATTCTATCAGTGAACCAAGAAGAATTAAACTATTCTATCGGTGTTGGGAGCAACATA
TTGGGGGAGAAGAACGCTCTAGTCTATGCAAGATGCTTACCAAGATCTGAGGGGATTATGGCTCAAAACGTC
GTGAAGTTATTCAAAAGCTCATGGAAAAGCTCTAGTTAAATGCAATTGGTCATTATCAAGAGATTCTAAAGCTTGG
TGACAATATGCAACCAATTGATTGCTGCAAAAGTATCCCTAACCTGCTAGGCTATAAAGATTCAATTAAACCATIG
AAACTCAGCAGTTGCAAGGGTGAAGGAGAGATACTCAAGGTGTTCTACAGATTGTTGAGAGCACATATGTTGAAAT
GCAAAATCTACTCCGACGTTCTTGGACAAGAAGAAGTACTCAGCTTGAACCAATGCGGTTTATTGAAAGATTATTG
GCAGAGCACCTGTTGGTGTGAGAAGATAATTAA

4256.2 (SEQ. ID. NO. 359)

ATGAAAAAAACAAGCTATGTCATTATTGCTCTCACCTCCTTCTATTGCTTTTCTCCACAGCTTGTGGAATACTT
GATTTGACTGGTCTATCTTTCGACGATGTCGAAAAAACAGAAAAATTGCTTTATTGTTGGTTTCAGCATGTCAT
ACCTGTCTCTAGCCCTGTTGGCAGGGATCGAAGAGCTTCTCAAGAAAATGCAAGGCTAATCTCAAGCGTTATTAGC
AGGGCTAACAAAGCTAAAATCAGAGCTTGTCTAGGAGAAGGAAATCATCGAGAACCGAAGCGAATTGCTCG
GATTGACGATACAGTCAGTCAAGGAGTTGTCAGAGTGTACAGCTATTAGAAAACAGCCAGAAGGATTGCGGTTTGCTCTG
TAGAGAAAAGATGCAAGACCCAGTTGAGCAGAGCTGATAGAAGGGATCAAATTCTTAAAGAGCTGAGGACAAGAGTG
CATTTGCGACCAAGTGAAGTGGAGCAGAAGAGCTGATAGAAGGGATCAAATTCTTAAAGAGCTGAGGACAAGAGTG
ATCTTAGGGTGTAGTCTCAAGCAGAAATTGACGAAATTGCTTAAGAAAATCGAGGAGCATATCTCCGATATCCTGCAAGAGTT
GATTAGCAATTCCCTCCGATCTGCTAGGAGCATCTGCTAGATGTCACCTCTATCAGACAGATGTTGAATTGCAACTGAGG
TGGTGGACAATGGGATTGGTTCCAGTTAGGGAGCTTAGAGCAGCTGAGTTATGGACTGCGAAATATCAAGGAGCGGGTTGA
AGATATGGCTGGAACAGTTCAACTCTGACAGCTCCAAGCAAGGGCTGGCGTTGATATCCGTATTCCCTGTTAGATAAG
GAATGA

4263.1 (SEQ. ID. NO. 360)

ATGATTGTTCCATTATTCTCAAGGATTGCTGGCTATTCTAGGTCGGGAATCTTATGACATTAGGATTTAAACTT
CCAGATATGACGACAGAACGGTTCCCTCCCTCTGGGGAGCTGTTGCTGTCATTGATAACCAAGGCGTAACCCATT
AGCGACACTTGTGCTGAGGAGCAGGTTGTTGGCTGGAATGGCAGCAGGGCTTCTTATACAAAAGGAAAGATCCAACC
TTGCTCTCAGGGATTGGTGTGACTCTTGTACTCAATGCTTGTATTGGACGTGCGAATTAGGCCTGCTTGG

ACCAAGCAAATTCAAGGATGTTTGCCTTGATCGGATTGAACTCAACTCTGACAGGTCTCATTTGTGAGTATTGGTATT
GCTCTCATGCTCTTTCTGGACACTAAACTCGGACAACGCTATATTGCTACAGGGATAATCCTGATATGGCTAGAAGTT
CGGGATTCACTGGACGCATGGAGCTCATGGGCTTGGCTTATCAAATGGTGTGATTGCCCTGAGGTGCCCTCATTGCTC
AGCAAGAAGGTTATGCCATGTCGAGGGATCGGGTTATCGTTGTTGGCTGCAAGTTGATTATGGAGAAGTTAT
TTCAAGAGTTGAGCTTGGCAGAGCGTTGGTACTATCGTTGAGGTCTATCGCTTATCAATTAGTGGCAGTT
CGCACTGGCTTAATACAAGTACCTCGTTATACAGTGCCTGATTAGCAGTCTGCCTCATGATCCAACATTTAACAGCA
AACATCTGAAAGGAGCCAAGTTAACAGCAATGA

4346.1 (SEQ. ID. NO. 361)

ATGAAAAAAATGAAAGTTGGTCACTGACTTGCAACGGGAGTTGCTTACTACACTTGTGCTGCTGGAGGTTCAAA
TTCTACAGTCTCTTCATCTGAAGAAAAAGCTGATAAAAGTCAGAATTAGTTATCTATTGAACTCAGTCTCAATGGTC
GTGGTGTATTGCTGAGAGAATAATGCAAGTGGTTAATATAAAAATGGTTGATATCGCTGGCCTCAATTAGCAGA
CCGTGTATTGCTGAGAGAATAATGCAAGTGGTTAATATAAAAATGGTTGATATCGCTGGCCTCAATTAGCAGA
AAAAATTACTAGTACAGTACAAGCTTAAATGGTAGATAAAATTGATCAATCTTATCAGATAAAGATAATTATAATCCT
GTGATTGTTCAACCATTAGTTAAATTGGGCGCTGATGAAAAGAAATGCTAAAGATTGGACTGAATTAGGTAGTAAGT
ATAAGGTTAATTCATTTCTGCTTCAAGGAGGTACAGGACGGCAATTCTAGCAAGTATCTAGTTGACACCTTGAT
GATAAGGTTAATAGGTGTTCCGAAAAAGGGTGGAGTAGCAAAGAATATTGAAAATGCAACACTCTTCAAAAG
GGAGAAAGTTCAATTGTTAGAGAAGATGTTAGACAAAGAAGATCTGAGATTGGTGTACCATTTGAACTGAACAAACTATGGTTTA
AGCACTAGAAAAACAAAGCTTAGCTAAAGAATTATTGATTGGTTGGTCAATCAGAAATTCAAGTAGAATATAGTAAGA
ACTTTGGATCTATCCCTGCAAAATAAGATGCCCTCAAAGATCTACCTGAAGAGATACGAAGAAATTGGTGTCAAGTGAAC
ACAAAATATTGACTGGGAAAGCTGTTGAAAGCATTGGATGAATGGTAGAAAAGCTGAATTAGAATACGTACAATAA

4346.2 (SEQ. ID. NO. 362)

ATGATTAATTTGATAATATTCAAATTAAATATGGTATTGCAATTGATAATCTGAATTAGATATACTGAAGGGGA
ATTTTTACATTCTGGCCTCAGGATGTTAAATCAACTACTTTGAGAGCATTGTTAGGTTCTAGATCCATCATCAGG
AAGTATTGAAAGTTAATGCAAGAGATGCACTATTGGAAACCTGAAAAGCGTGGATTGGTATTGATTTCAATCTTATGCGC
TATTCCAACATGACTTTGATAATATTGCAATTGGTTAAAAGTTAAGAAGGTAGCTCAGGATGTTATTAAAGCTAA
GTATCAGCAGTGGCAGCAAAATTAAAGATCTGATCACAGTTACAGCGTAATGTTAGCAGATTATCTGGGGTCAACAC
AAAGGTTAGCATTGGCTCGTCTGGTCTGAACTTAAATTCTGTTAGATGAACCATGTCACCTTGACGCAAA
TTACGTGTAGATTGAGAAAAGAGTTGAAAAGACTTCAAAAGAGTTAGGTATTACTACTTTATATGTTACTCATGATCAAG
AGGAAGCCTGACTTTGATGAGAATTGCAAGTCTTAACAATGGATACATCGAACAGTCCGTAACCCAGTAGAGATTAA
TCATAATTCTCAAAGTGAATTGTTGATTGTTATTGGAGATAATTGTTGAGCTGACGTTACGTCACGAAACTATTAT
TGAAAATACAGCGTTCTTCAATTAGGATAAAAAGGATAATTGCAATTGTTGAGGTTACAATTCAACTATACAATAAAGTTCTGAAAGTC
AGATTCTTAATGTAACAAGTATTGATGTTGAGTTCTGGAGTTACAATTCAACTATACAATAAAGTTCTGAAAGTC
GACGTTCTGCAATTITAA

4346.3 (SEQ. ID. NO. 363)

ATGCGTCATAAATTAAAGATTGGCTTATTGTTAGGGTTAATCTGGTCTTAGTAACATTATTATTCATCAAAC
TTTGATCTAGTAGTGAATGTTAGTAAAGGAGGAGAATTTCCTGATGCTGATCATCGTCTAAATCTCAGAGGGC
ACTTCAGAGTATTATGAAACAGTTAAGTTAGCATTTCACCTATTACAGTTAATGCTGTAGGTATTCTTGTGTTCTATT
TACAGAGTACTTGTGATATTAAAGGTGCTAAATTAAATTAGGTTATATGACCTCTTAAATTGAGGAGTGGTTTTAG
CGACTGGATATAATTGCTATGGCTTATGGATTGATTACAAAATTTCACAAATGTTATCCCTCTTAAACCCCTAATC
GGTTTATTGGGATGGTCAGTCTTATTGACATTTCAGGAACCTGCTAATCATATTGTTAAACAAATACAATTIC
GAAGCGTTGACTATCACACTATTGAGGCTGCTGAAATATGGAGCAAAACCAATTACTGTTCCGAAAAGTAGTTTAC
AACCTTAATCCAACCTATTGCACTTACTATTATGGTTTCTAGTGGTTATCTGAGTAGCAGCACCCATGATTGTTGG
TGGTAAAGAATTCTAAACTATAAACCAATGATTACATTGCAAGGGATGGGAAATTCTGTTGATTAGCTGCCCTACTTG
CAATTATTAGGTATTGCAACTACAATTGCTTACTATCATGAATAAGATAGAAAAGGTGAAATTATTTCTATCTCT
AAGACTAAAGCGCTTAAACAAAATTGCGTCAAGCCTGGAATATCATTGTCACATTGTTAGCATATGGATTGT
TCACAGTTTCATGCTTCACTAATTATTAGTATTATCACTATTACAGATCCAGTTGCAATTCAAACAGTAACCTAACAT
TATCAAACATTACTTAAAGAAAATTATGCTTATTCTTGTAGTAATAGTGGCGCATTCTCTCCATTGGTCACTTATTATC
TATTATTGCTGCAACACAGCAACAATTCTCGCAGTTGATTGCTGTGTTGCAAAAACATAATCTGTTTGTATTCT
ATTGAAATATTGCTACTTCTGCTTACTACCAAGTACACTTTACGAGTAAGTTATTACTTTAATCAGCCACA
ATTCTTGTCTTGAATCAGATTGGTAGGTATTGGTAATTCTACTTATTGCAATTAGTTGAAAATTCCATTCTTAT
AGAATGGTAGCTATTATTAGTGTGATGAGATGGAAGATGCAAGAAGTATGGGTGCTTCACCTTTTATAC
TATGATGAAGGTTATCATTCCATTATTACCGGTTCTCTCTGTATTGTTAAACTTAACTCTTATTAACGTACTTC
GACTTATCTGATTCCATTACCATCCCTAGCTAACCAATTAGGTATTACGATTGATCTGCAGGTGATGAAACAGCAACATC
TAATGCACAAGCTCTGGTATTGTTATACAATTGTTGATGATTATTCTGGAACGGTATTACTTCACACAAAGACCGG
GGCGTAAAGTAAGGAAATAA

Table 2**(SEQ. ID. NO. 1)**

MEEVLTLDCFLIDRTKIEANANKYSFWWKKTTEKFSAKLQEIQVYFQEEITPLLIKYAMFDKKQKRGYKESAKNLANWHYNDKE
DSTHDPDGWYYRFHHTKYQKTQTFQQEIKVYYADEPESAPQKGLYMNERYQNLKAKECQALLSPQRQIQAQRKIDVEPVFGQI
KASLGYKRCNLRGKRQVRIDMGLVLMANNLLKYSKMKZ

(SEQ. ID. NO. 2)

MGKGHWNRKRVYSIRKFAVGACSVIMGTCAVLLGGNIAGESVVYADETLITHTAEKPEEKMIVEEKADKALETKNIVERTEQSE
PSSTEAIASEKEDAEAVTPKEEKVSAKPEEKAPRIESQASNQEPLKEDAKAVTNEEVNQMIEDRKFVDFQNWNWFKLNANSKEAIK
PDADVSTWKLLDLPYDWSIFNDHESPAQNEGGQLNGGEAWYRKTFLDEKLKVNRLTFDGVYMDSQVYVNGQLVGHP
NGYNQPSYDITKYLQKDGRENVIHAVAVNQPSRSWYSGSIYRDVTLQVTDKVHVCKNGTTILTPKLEEQHQGKVETHVTSKIV
NTDDKDHELVAEYQIVERGGAHTGLVRTASRTLKAHESTSLDAILEVERPKLWTVLNDKPALYELTRVYRDGQLVDAKKDLFG
YRYYHWTPNEGFSLNGERIKFHGVSLHHGDHALGAEENYKAERYRLKQMKEMGVNSIRTTHNPAEQLQIAELGLLVQEEAF
DTWYGGKKPYDYGRFFEKDATHPEARKGEKWSDFDLRTMVERGKNNPAIFMWSIGNEIGEANGDAHSLATVKRLVKVVKDV
TRYVTMAGDKFRGNGSGGHEKIADELDAVGPNYSEDATRTRGSYRPERELKHSNGPERNYE
QSODYGNDRVWGKTTATASWTFRDNAGYAGQFIWTGTDYIGEPTPWHNQNPVCKSYFGIVDTAGIPKHDFTYLSQSQWVSVK
KKPMVHLLPHWNWENKELASKVADSEKGIPVRAVSNASSVELFLNGKSLGLKTFNKQTSDFRTYQEGANANELEYLEWKVAYQ
PGTLEAIARDDESCKEIARDKITTAGPKAAVRLIKEDHAIAADGKDLTYIYYEIVDSQGNVPTANNLVRFQLHGQQQLVGVVDNGEQ
ASRERYKAQADGSWIRKAFNGKGVAVKSTEQAGKFTTAHSDLLKSNSQVTFTGKKEGQETKVLGTEVPKVQTIIGEAPEMPTT
VPFVYSDGSRAERPWTWSVDSVSKPGIVTVKGMAADGREVEARVEVIAKSELPPVVKRIAPNTDLSVDSVSVYVLIDGSVEEYEV
KWEIAEEDAKLAIPGSRIQATGYLEGQPIATLVEEGNPAPA VPTVTVGGEAVTGLTSQKPMQYRTLAYGAKLPEVTAASKN
AAVTVLQASAANGMASIFIQPKDGGPLQTYAIQFLEEAPKIAHSLQVEKADSLKDQTVKLSVRAYHQDGTQAVLPADKVTFS
TSGEVEAIRKGMLLEHHPGAVTLNAEYEGAKDQVELTIQANTEKKIAQSIRPVNVVTDLHQEPSPSLPATVTEYDKGFPKTHVKT
WQAPIKEKLDSYQTFEVLKGVEGIDLEARAKVSEGVISVEEVSVTTPIAEAPQLPESVRTYDSNGHVSAAVWDAIRPEQYAKE
GVFTVNGRLEGTQLTTKLHVRVSAQTEQGANISDQWTGSELPLAFASDSDNSPDPVSNVNDKLISYNNQ PANRWTNWNRNTNPEASV
GVLFGDGSILSKRSVNDLNSVGFHEDHGVPKSVIEYYVGKTVPTAPKNSPVGNDHVFNDSANWKPVTLKAPAQKLKAGEM
NHFSFDKVETYAVRIRMVKADNKRGSITESVQIFAKQVAAAKQGQTRIQVGDGKDLANFPNDLTDYLESVDGKVPVAVTASVNN
GLATVVPSPVREGEPEVPRVIAKAENGDIILGEYRLHFTKDKSLLSHKPVAAVKQARLLQVGQALELPTKVPVYFTGKDGYETKDLTV
WEEVPAENLTAKQFTVRGRVLGSNLVAEITVVTDLGETLSDNPVNDENSNQAFASATNDIKNSHDRVVDLNDGDHSENRR
WTNWSPTPSSNPEVSAVGIVFRENGKIVERTVTQKGVQFFADSGTDAPSKLVLERYVGPFEVPTYYSNYQADADHPFNNPENWE
AVPYRADKDIAGDEINVTFKAIKAKAMRWMERKADKSGVAMIEMTFLAPSELPQESTQSKILVDGKELADFAENRQDYQITYK
GQRPKVSVEENNQAVASTVVDGEDSPVVLVRLVSESQKQVKEYRIHLTKEKPVSEKTVAAVQEDLPKIEFVEKDLAYKTVEKKDS
TLYLGETRVEQEGKVGKERIFTAINPDGSKEEKLREVVEVPTDRIVLVTGPVQAQEAKKPQVSEKADTKPIDSSEASQTNKAQPLST
GSAASQAAVAAGLTLGLSAGLVVTKGKEDZ

(SEQ. ID. NO. 3)

MKIMKKKYWTLAILFFCLFNNSVTAQEIPKNLDGNITHQTSESFSDEKQVDYSNKNQEEVDQNKFRIQIDKTELFTVTTDKHLEK
NCCKLEPQINNDIVNSESNNLLGEDNLDNPKENVSHLDNRGGNIEHDKNLESSIVRKYEWIDKVTGGGESYKLYSKNSKV
SIAIDSGVDLQNTGLLKLNLSNHSKNYVPNKGYLKEEGEEGIIQDRLGHGTAVVAQIVGDDNINGVNPHVNINVYRIFGKSSA
SPDWIVKAIFDAVDDGNDIINLSTGQYLMIDGEYEDTNDFTFLKYKKAIDYANQKGVIIAALGNDLSVNSNQSDLKLIRRKK
VRKPGLVVDVPSYFSSTISVGIDRLGNLSDFSNKGDSDAIYAPAGSTLSSELGLNNFINAEKYKEDWIFSATLGGTYLYGNSFA
APKVGAIAMIIDKYKLKDQPYNMVFVKKFWKKHYQZ

(SEQ. ID. NO. 4)

MKKTWKVFLTLVTALVAVVLVACQGQGTASKDNKEAELKKVDFILDWTPNTNHTGLYVAKEKGYFKEAGVDVDLKLPPPEESSSD
LIVINGKAPFAVYFQDYMAMKKLEKGAGITA VAAIVEHNTSGIISRKSDNVSSPKDLVGKKYGTWNDPTELAMLKTLVESQGGDFEK
VEKVPNNDSNSITPIANGVFTA WIIYGDGILAKSQGV DANFMYLKDVKEFDYSPVIIANNNDYLKDNEKEARKVIQAIKKGY
QYAMEHPEEAADILIKNAPELKEKRDVFVIESQKYLKEYASDKEKWGQFDAARWNANFYKWDKENGILKEDLTDKGFTNEFVKZ

(SEQ. ID. NO. 5)

MKRTWRNSFTNLNTPFMIGNIEIPNRTV LAPMAGVTNSAFRTIAKELGAGLVMEMVSDKGIQYNNNEKTLHMLHIDEGENPVS
QLFGSDEDSLARAEEFIQENTKTDIVDINMGCVPNKIVKNEAGAMWLKDPPDKIYI SIIKVKQSVDIPLTVKMRTGWADPSLAVENA
LAAEAAGVSALAMHGRTRQMYTGHADLETLYKVAQALT KIPFIANGDIRTVQEAQKRIEVEVGADAVMIGRAAMGNPYLFNQIN
HYFETGEILPDLTfedKMKIAYEHLKRLINLKGENVA VREFRGLAPHYLRGTSGAAKLRAISQASTLAEIETLLQLEKAZ

(SEQ. ID. NO. 6)

MKPNPKLLTSFLRSFAILGGVGLVIIHAIYLTFPFYYIQLEGEKFNESARVFT EYLKTKTSDEIPSLLQSYSKSLTISAHLKRDIVDKR
LPLVHDLIDIKDGKLSNYIVMLDMSVSTADGKQTVQFVHGVVYKEAKNILLLYPFTFLVIAFSFVFSYFYTKRLLNPLFYISEV
TSKMQDLDNNIRFDESRKDEVGEVKGQINGMYEHLLKVIYELESNEQIVKLQNKVSVFVRGASHELKTPASLRRILENMQHNIG
DYKDHPKVIASKINKIDQMSHLLVELESSKFQEWTECRETTVKPVLVDILSRYQELAHSIGVTIENQLTDAVVMSLRALDKVLT
NLISNAIKYSDKNGRVIISSEQDGYLSIKNTCAPLSDQELEHLDIFYHSQIVTDKDESSGLGLYIVNNILESQYQMDYFLPYEHGMF
ISLZ

(SEQ. ID. NO. 7)

MYLGDLMEKAECGQFSILSFLLQESQTTVKAVMEETGFSKATLTKYVTLNDKALDGLLELAIHSEDENRLSICAATKGRDIRSLF
LESAVKYQILVYLLYHQQLAHQLAQLQELVISEATLGRHLAGLNLQILSEFDLSIQNQGRWRGPEHQIHYFYFCLFRKVWSSQEWEGHM
QKPERKQEIANLEEICGASLSAGQKLDLVLWAHISQQRRLRNACQFQVIEEKMRGYFDNFYRLRLRKVPSFFAGQHPLGVEDGE
MMIFFSFLSHRILPLHMTMEYILGFGQLADLLTQLIQEMKKEELLGDYTEDHVTYELSQLCAQVYLYKGYILQDRYKYQLENRHP
YLLMEHDFKETAEEIFHALPAFQQGTLDKKILWEWLQLIEYMAENGQHMRIGLDDLSGFLVFSRMAAILKRYLEYNRFITIEAY
DPSRHYDLLVTNNPIHKKEQTPVYLYKNDLDMEDLVAIRQLFTZ

(SEQ. ID. NO. 8)

MEFSKKTRELSEKKMQERTLDLLIIGGGITGAGVALQAAASGLETGLIEMQDFAEGTSSRSTKLVHGLRLYKQFDVEVVS DTVSE
RAVVQQIAPHIPKSDPMPLLIVYDEDGATFSLFRLKVAMDLYDLLAGVSNTPAANKVLSKDKVLERQPNLKKEGLVGGGVYLDFR

NNDARLVIENIKRANQDGALIANHVKAEGFLFDESGKITGVVARDLLTDQVFEIKARLVINTTGPWSDKVRNLSNKGTQFSQMRPT
KGVLVVDSSKIKVSPQVYFDTGLGDRMVFLPRENKTYFGTTDVTGDLHPKVTQEDVDYLLGIVNNRFPESENITIDIES
WAGLRPLAGNSASDYNNGNNGTISDFDNLIATVESYLSKEKTREDVESAVSKLESSTSEKHLDPSAVRGSSLRDDNGLTLA
GGKITDYRKMAEGAMERVVDLKAEDRSFKLINSKTPVSGGELNPANVDSEIEAFAGLGVSRLDSKEAHYLANLYGSNAPKV
FALAHSLEQAPGLSLADTLSLHYAMRNELTSPVDFLLRTNHMLFMRDSDLSIVEPILDEMGRFYDWTEEEKATYRADVEAALA
NNDLAELKNZ

(SEQ. ID. NO. 9)

MMNELFGEFLGTLILILLGNGVVAGVVLPKTSNSSGWIVITMGWIAVAVAVFSGKLSPAYLNPAVTIGVALKGGLPWASVLP
YILAQFAGAMLGQILVWLQFKPHYEAEEENAGNILATFSTGPAIKDTSNLISEILGTVLVTIFALGLYDFQAGIGTFAVGTLIVGIG
LSLGGTTGYALNPARDLGPIMHSILPIPKNKGDGDWSYAWIPVVGVPVGAALAVLFSLFZ

(SEQ. ID. NO. 10)

MTKKKIERISVIHREKILWLKWYFMRDKEQPKYSVLERKMFDAAKNQDMAVYQKYATIKQITDIRVQTSEADILEAVKEVYVYNH
MNVIGACQRILFISQSPAYDKLNKFNFNIYSDLYFSVVPKMGVYHEMVGIZ

(SEQ. ID. NO. 11)

MKNSNEAEMKLLYDRTSLTEILTREAEELVAAGKRVFYIAPNSLSFEKERAVLEYLSQQSFSITVTRFAQMARYLVNLDPAKT
TLDDIGLGLAFYKCLAEQDKDLRVYGAIKQDPQLIQQLIELYHEMTKSQMSFLDLENLTDEDKRADLLIFEKVTAYLNQQLAQ
ESQLSHLIEAIENDKVSSDFNQIALVIDGFRFSAEERVVVDLLHGKVEIVIGAYASKKAYTSPFSEGNNLYQASVKFLHHASKYQT
PAQDCSQTKEKMDSFDAKSLLESSYDFSELALDVEKDRENQIWSCLTQKEELELVARSIRQLHENSDSLQYKHFRIILGDVAS
YQLSLKTIFDQYQIPFLGRLSEAMAHPLTQFVESILAKRYFRQEDLNLRTDLYTDLQSDDIAFEQYIRYLGINGLPAFQQTFT
KSHHGKFNLERLNVLRLRAPLETFLASRKQKAELLQKWSVFLKEGAVTKQLQDLTTLEAVEQERQAEVWKAFCHVLEQF
VFAGSQVSLLEDFLALLHSGMSLSQYRTIPATVDTVVLVQSYDLIAPLTDVFVYAIQLTQDNLPKISQNTSLLTDEERQNLNQATEEGV
QLLIASSENLKKNRYTMSLVSARKQLFLSAPSLFNESESKESEYLQELIHFGFRRREKRMNHKGSKEDMGSYHSLLSLVAYH
QQGEMSDTEQDQDFVVKVLSRVIGKLLDQQGLENPAPTSPSSKTLAKDTLQALYPAKQEFYLSLTSGLTEFYRNEYSYFLRYVGLQ
EELRLHPDASHGNFLHRIFERALQPNEDSFQDRLQEAQIQTQSEREFEATYQESLEAQFTKEVLLDVARTTGHILRHNPAAETIKEE
ANFGGKDQAFIQLDNGRSVFRGKVDRIDRLKANGAIGVVDYKSSLTQFQPHFFNGLNSQLPTYLAALKREGEQNFFGAMYLEM
AEPVQSLMAVSLAGAVVEASKSMKYQGLFLEKESSYLGEFYVNKNKANQLTDEEFQLLDDNAYLYKAAEKILAGRFAINPYT
ENGRSIAPVQQHQAITGFEANYHLGQARFLEKLDLADGKRLVGEKLKQAWLEKIREELNRZ

(SEQ. ID. NO. 12)

MKLIPFLSEEEIQKLQEAEANSSKEQKKTAEQIEAITYSTAQNILVSASAGSGKTFVMAERILDQLARGVEISQLFISTFTVKAATELKE
RLEKKISKKIQETDDVDLKQHHLGRQLADLPNAIGTMDSTQFLGKHGYLLDIAPNFRILQNCSEQLILENEVFHEVFEAHYQGK
QKETFSHLLKNFAGRGRKDERGLRQQVYKIYDFLQSTSNSPKWLSESFLKGFEKADFTSEKEKLTEQIKQALWDLESFFRYHLDND
AKEFAKAAYLENVQLLDEIGSLNQESDSQAYQAVLARVVAISKEKNGRALTNASRKADLKPALDAYNEERKTQFAKLGQLSDQI
AILDYQERYHGDTWKLAQTFQSFMDFVEAYRQRKRQENAEFADISHYTIEILENFPQVRESYQERFHEVMVDEYQDTNHIQER
MLELLSNHGRNMVGDIKQSIYRFRQADPQIFNEKFQRYAQNPQEGRLIILKENFRSSSEVLSATNDVFERLMDQEVGEINYDNKH
QLVFANTKLTPNPDNKAALYDQDDTGEEEESQETKLTGEMRLVKEILKHQEKGVAFKEIALLTSSRSRNDQILLALSEYGIP
VKTDGEQNNYQSLLEVQVMLDTLVRVHNPLQDVALVLMKSPMFGFDEDELARSLSLQKAEDKVHENLYEKLVNAQKMASSQKG
LIHTALAEKLKQFDILASWRLYAKTHSLYDLIWKIYNDRFYYDVGALPNGPARQANLYALALRADQFEKSNFKGLSRFIRMID
QVLEAQHDLASVAVAPPDAVELMTIHKSGLEFPYVFLNMDQDFNKQDSMSEVLSRQNGLGVYIYAKMETGAVEDHYPKTIK
LSIPSITYRQNEEEQLASYSEQMRLLYVAMTRAEEKLYLVGKGSREKLESKEYPAAKGKLNNSNTRLQARNFQDWLWAISKVFT
KDKLNFSYRFIGEDQLTREAGELETKSPLQDSSQADNRQSDTKEALEMLKEVEYNTLHRAAIELPSVQTPSQIKKFYEPVMDME
GVEIAGQQGSVGGKISFDLPDFSTKEVTKGAEIGSATHELMQRIDLSQLTASLTETLKVQTSQVDRKINLDKILAFFDTVLGQ
EILANTDHLYREQPFSMLKRDQKSQEDFVVRGILDGYLLYENKIVLFDYKTDYDEPSQLVDRYRGQLALYEALSRAYSIENIEK
YLILLGKDEVQVVKVZ

(SEQ. ID. NO. 13)

MELARHAESLGVDIAITIPIYFRLPEYSVAKYWNDISSAAPNTDYVIYNIQPLAGVALTPSLYTEMLKNPRIVGVKNSSMPVQDIQ
TFVSLCGGEDHIVFNGPDEQFLGGRMLGARAGIGGTYGAMPFLKLNQLIADKDLTARELQYAINAIIGKLTSAHGNMYGVIKEV
LKINEGLNIGSVRSPLTPVTEEDRPVVEAAAALIRETKERFLZ

(SEQ. ID. NO. 14)

MYTKCLREKLVLFLKIFFPILYQFANYSASFVDTAMTQGQYNTMDLAGVSMATSIWNPFFTFLTGIVSALVPIIGHHLGRGKKEEV
ASDFYQFIYLALGLSVVLLGMVLFLAPIIHNHIGLEAAVAAVAVRYLWFLSIGIIPLLLFSVIRSLLDGLTKLSMYLMLLLPLNSG
FNYLLIYGAFCVPELGGAGAGLGTSLAYWVLLGISVVLFLQKQELKALHLEKRIPLNMDKIKEGVRGLPIGGTVFAEVAIFSVVGL
IMAKFSPLIIASHQSAMNFSSLMYAFPMISSAMAIVVSYSYEVGAKRFDAAKTYIGLGRWTALIFAATLTFLYIFRGNVASLYGNDP
KFIDLTVRFLTYSLLFQLADTFAPAQLQGILRGYKDTVIFPFLGGLGYWGVAIPVYAI

(SEQ. ID. NO. 15)

MSTLAKIEALLFVAGEDGIRVRQLAELLSLPPGIQOSLGLKLAQKYEKDPDSSLALIETSGAYRLVTPQFAEILKEYSKAPINQSL
RAALETLSIIAYKQPITRIEIDAIRGVNSSGALAKLQAFDLIKEGKKEVLGRPNLYVTTDYFLDMGINHLEELPVIDELEIQAQESQ
LFGERIEEEDENQZ

(SEQ. ID. NO. 16)

MDTMISRFFRHLFEALKSLRNGWMTVAAVSSVMITLTLVAIFASVIFNTAKLATDIENNVRVVVYIRKDVEDNSQTIEKEGQTVT
NNDYHKVYDSLKNMSTVKSFTSSKEEQYEKLTEIMGDNWKIFEGDANPLYDAYIVEANTPNDVKTIAEDAKKIEGVSEVQDGG
ANTERLFKLASFIRVWGLGIAALLIFIAVFLISNTIRIHSRSREIQIMRLVGAKNSYIRGPFLLEGAFICLLGAIAPSVLVIVYQIVYQS
VNKSLVGQNLSMISPDLFSPLMIALLFVIGVFIGSLGSGISMRRFLKIZ

(SEQ. ID. NO. 17)

MKKVRFIFALLFFLASPEGAMASDGWTQGKQYLKEDGSQAANEWVFDTHYQSWFYIKADANYAENEWLKGQGDDYFYLKSGG
YMAKSEWVEDKGAFYILDQDGKMRNAWVGTSYVGATGAKVIEDWVYDSQYDAWFYIKADGQHAKEWLQIKGKDYYFKS
GGYLLTSQWINQAYVNASGAKVQQGLFDKQYQSWFYIKENGNYADKEWIFENGHYYLKSGGYMAANEWIWDKESWFYLK
FDGKMAEKEWVYDHSQAWYYFKSGGYMTANEWIWDKESWFYLKSDGKIAEKEWVYDHSQAWYYFKSGGYMTANEWIWD
KESWFYLKSDGKIAEKEWVYDHSQAWYYFKSGGYMAKNETVDGYQLGSDGKWLLGKTTNENAAYYQVVPVTANVYSDGE
KLSYISQGSVVWLDKDRKSDDKRALATISLGSYMKTEDLQDALSKDFPQYYESDGHRFYHYVAQNASIPVASHLSDMEVGKKY
YSADGLHFDFGKLENPLFKDLTEATNSAEELDKVSFLINNSLNENKGATFKEAEHEYHINALYLLAHSALESNWGRSKIADK
KNNFFGITAYDTTPYLSAKTFDDVDKGILGATKWIKENYIDRGRFLGNKASGMNVEYASDPYWGEKIASVMMKINEKLGGKDZ

(SEQ. ID. NO. 18)

MKKVLQKYWAFAVVIPLLLQAIFYVPMFQGAFYSFTNWTGLTNYKFVGLNNFKLLFMDPKFMNAIGFTAIIIAIAMVVEIAL
GIFIARVLSNKIKGQTFFRAWFFFPAVLSGLTVALFKQVFNYGLPAIGNALHIEFFQTSLGKWTGAIFAAVFVLLWQGVAMPIIFL
AGLQSIPIITEAARIDGATSKQVFWMIELPYLLPSVSMVFLALKGGLTAFDQVFAMTGGPNNATTSLGLLVNYAFKNNQFGY
ANAIAMILFFLIVVISIQLRVSKKFEIZ

(SEQ. ID. NO. 19)

MMKQDERKALIGKYILLGSLVILVPLLATLFSSFKPTKDIDNFQGPTNFTWDNFSRLLADCGIGGYYWNSVITVLSLLAVMIFP
MAAYSIARNMSKRKAFTIMYTLILGIFVPFQVIMIPITVMMMSKGLANTFGLILLYLTIAIPQTLFLYVGYIKISIPESLDEAAEIDGA
NQFTTYFRIIIPMMKPMHATTMIINALWFWNDFMLPLLVNRDSKMWTPLFQNYAGQYFNDYGPSFASYVVGIIISITIVLFFQ
RHIISGMSNGAVKZ

(SEQ. ID. NO. 20)

MKSILQKMGEHPMLLFLSYSTVISILAQNWMGLVASVGMFLFTIFLHYQSIILSHKFRLILQFVLFGSVLSAAFASLEHFQIVKKF
NYAFLSPNMQVWHQNRAEVTFNPNEYGIICFCIMIAFYLFTTCKLNWLKVFCVIAGFVNLFGLNFTQRNTAFPAIIAGAIYLFTT
KNWKAFLWSIGVFAIGLSFLFSSDLGVRMGTLDSSMEERISIWDAAGMALFKQNPFWGEGPLTYMSYPRIHAPYHEAHSLYIDTI
LSYGIVGTILLVLSSVAPVRLMMMDMSQESGKRPIGLYLSFLTVVAHGIFDLALFWIQSGFIFLLVMCSIPLEHRMLVSDMTDZ

(SEQ. ID. NO. 21)

MSKMDVQKIAIPMMKVFVNMRGIIALKDGLAILPLTVVGSLFLIMGQLPFEGLNKSIAVFGANWTEPMQVYSGTFAIMGLISCFS
IAYSYAKNSGVEALPAGVLSVS AFFILLRSSYIPKQGEAIGDAISKVWFGGGQIIGAIIGLUVGSIYTFIIRKRVIKMPEQVPOAIAKQ
FEAMIPAFVFLSSMIVYILAKSLTNGTFIEMIYSAIQVPLQGLTGSYGAIGIAFFISFLWWFGVHGQS VVNGVTALLSNLDAN
KAMLASANLSLENGAHIVTQQFLDSFLILSGSGITFGLVVAMILFAAKSKQYQALGKVAAPFAIFNVNEPVVFGPIVMNPVMFVPFI
LVPVLAIVVYGAITGFMQPFSGVTLWPWSTPAISGFLVGGWQGVITQLVILAMSTLVYFPFFKQVQDRLAYQNEIKQSZ

(SEQ. ID. NO. 22)

MKKKDLVDQLVSEIETGVRTLGIYGHGASGKSTFAQELYQALDSTTVNLLETDPYITSGRHLVVPKDAPNQKVTASLPVAHELES
LQRDILACRRVWMSZ

(SEQ. ID. NO. 23)

MKKRYLVLTALLALSLAACSQEKTKNEDGETKTEQTAADGTVGSKSQGAAQKKAEVVNKGDDYYSIQGKYDEIIVANKHYPLSK
DYNPGENPTAKAELVKLIKAMQEAGFPISDHYSGFRSYETQTKLYQDYYVQNDGKAAADRYSARPGYSEHTGLAFDVITDGDL
VTEEKAAQWLLDHAADYGFVVRYLKGKEETGYMAEEWHLRYVGKEAKEIAASGLSLEYYGFEGGDYVDZ

(SEQ. ID. NO. 24)

MREPDLNHFLKKGYFKKHAKAVLALSGGLDSMFLFKVLSTYQKELEIELILAHVNHKQRIESDWEEKELRKLAEEAELPIYISNFS
GEFSEARARNFRYDFFQEVMKKTGATALVTAHHADDQVETIFMRLIRGTRLRLSGIKEKQVVGIEIEIIRPFLHFQKKDFPSIFHFED
TSNQENHYFRNRIRNSYLPLEKENPRFRDAILGIGNEILDYDLAIELSNINVEDLQLQFLSYSESTQRVLLQTYLNRFPDNLNTKA
QFAEVQQILSKSKSQQYRHPKNGYELIKEYQQFQICKISPQADEKEDELVLYQDNQVAYQGQLFSFGLPLEGELIQQIPVSRETSIHIRH
RKTGDVLIKNGHRKKLRLFIDLKIPMEKRNSALIIEQFGEIVSILGIATTNLSKKTNDIMNTVLYIEKIDRZ

(SEQ. ID. NO. 25)

MRKFLIILLLPSFLTISKVVSTEKEVYVTSKEIYYLSQSDFGIYFREKLSSPMVYGEVPVYANEDLUVVESGKLTPKTSFQITEWRLNK
QGIPVFKLSNHFIAADKFLYDQSEVPTIKVWLESDFKLVLNPSYDLKEVKSSLAYSQVSIDKTMFVEGREFLHIDQAGWVAK
ESTSEEDNRMSKVKQEMLSKEYQKDFSIYVQKLTGKEAGINQDEKMYAASVLKLSLYYYTQEKENIGLYQLDTTVKYVSAVNDF
PGSYKPEGGSGSLPKKEDNEKEYSLKDLITKVSKESEDNVAHNLLQNSQDAFTKSKMSAIMGDDWDPEKLISSKMGKFMEAI
YQNQGFVLESLTQDFDSQRIAKGVSVKVAHKICDAEFKHDGVVYADSPFILSIFTKNSDYDTISKIAKDVYEVLKZ

(SEQ. ID. NO. 26)

MKKQNNGLIKNPFLWLLFIFFLVTGFQYFYSGNNSGGSQQINYTELVQEITDGNVKELTYQPNGSVIEVSGVYKNPKTSKEETGIQF
FTPSVTKVEKFTSTILPADTTVSELQKLATDHKAETVVKHESSGIWINLLVSIVPGILFFFLSMMGNMGGGNRNPMSGRSKA
KAANKEDIKVRFSVDVAGAEEEKQELVEVVEFLKDPKRTKLGARIAGVILLEGPPGTGKTLAKAVAAGEAGVPPFSISGSDFEVEMF
VVGASRVRSLFEDAKKAAPAIIFIDEIDAVGRQRGVGLGGGNDEREQTLNLQLLIEMDGFEKGNEGIVIAATNRSDVLDPALLRGR
FDRKVLVGRPDVKGREALKVHAKNPKLAEDVDLKLVAQQTPGVGADLENVLNEAALVAARRNKSIIASDIDEAEDRVIAGPS
KKDKTVSQKERELVAYHEAGHTIVGLVLSNARVHKVTIVPRGRAGGYMIALPKEDQMLSKEDMKEQLAGLMGGRVAEEIIFN
VQTTGASNDFEQATQMARAMVTEYGMSEKLGPVQYEGNHAMLGAQSPQKSISEQTAYEIDEEVRSLLNEARNKAAEIIQSNSRETH
Kliaeallkyetldstqikalystgkmppeaveeeshalsydevkskmndeckz

(SEQ. ID. NO. 27)

MKRSSLVVRMVISIFLVFLILLALVGTFYYQSSSSAIEATIEGNSQTTISQTSIFIQSYIKKLETTSTGLTQQTDVLAYAENPSQDKVEG
IRDLFLTILKSDKDLKTVLVTKSGQVISTDDSVQMKTSSDMAEDWYQKAIHQGAMPVLTPARKSDSQWVISVTQELVDAKGA
NLGVLRLDISYETLEAYLNQLQLQGQQGAFIENHENHEFVYHPQHTVYSSSSKMEAMKPYIDTQQGTYTPGHKSYVSQEKIAGTDWTV
LGVSSLEKLDQVRSQQLWTLGASVTSLLVCLCLVWFSLKRWIAPLKDLRETMLIEASGAQNLRAKEVGAYELREVTRQFNAMLD
QIDQLMVAIRSQEETTRQYQLQALSSQINPHFLYNTLDIWIWMAEFHDSQRVVQVTKSLATYFRLALNQGKDLICLSDIEINHVRQYL

FIQKQRGYDKLEYEINENVAFDNLVLPKLVLQPLVENALYHGIKEKGQGHIKLSVQKQDSGLVIRIEDDGVGQDAGDSSSQLK
RGGVGLQNVQDQLRLKHFGANYHMKIDSRPQKGTKVEIYINRIETSZ

(SEQ. ID. NO. 28)

MKRSSLVVRMVISIFLVFLILLALVGTFYQQSSSAIEATIEGNSQTTISQTSHFIQSYYIKKLETTSTGLTQZTDVLA耶AENPSQDKVEG
IRDLFLTILKSDKDLKTVLVLTKGQVISTDDSVQMKTSSDMAEDWYQKAIHQGAMPVLTPARKSDSQWVSVTQELVDAGKA
NLGVLRLDISIYETLEAYLNQLQLGQQGFAFINENHEFVYHPQHTVYSSSKMEAMKPYIDTGGYTPGHKSYSVSEKIAQTDWTV
LGVSSLEKLDQRSSQLLWTLLGAWSPLLVCCLVWFSLKRWAIPKLDRRETMLIEASQAQNRLAKEVGA耶YELREVTRQFNAMLD
QDQLMVAIRSQEETTRQYQLQALSSQINPHFLYNTLDTIWMAEFHDSQRVVQVTKSLATYFRLALNQGKDLCILSDEINHVRQYL
FIQKQRGYDKLEYEINENVAFDNLVLPKVLQPLVENALYHGIKEGQGHIKLSVQKQDSGLVIRIEDDGVGFQDAGDSSSQLK
RGGVGLQNVQDQLKLHFGANYHMKIDSRPQKGTKVEIYINRIETSZ

(SEQ. ID. NO. 29)

MFFKLLREALKVQVRSKILFTIFLVFRIGTSITVPGVNANSLNALSGSFLNMLSLVSGNALKNFSIFALGVSPYITASIVQLLQ
MDILPKFVEWGKQGEVGRRLKNQATRYIALVLAFAQSIGITAGFNTLAGQLIKTALTPQVFTLTIQIILTGSMIVTWLGEQITDKG
YGNGVSMIIAGFIVSSIPEMIQGYIYDVYFVNVPSSRITSSIIFVIIIITVLLIYFTTYVQQAEYKIPQYTKVAQGAPSSSYPLKVNPG
VIPVFASSITAAPAAIQLQFLSATGHDWAVVRVAQEMLATTSPTEGAMYALLIILFTFFFYTFVQINPEKAETYKRVVPISEMFVLVK
VOKNICLNFFVVLLOLVPSSLVZ

(SEQ. ID. NO. 30)

MDIRQVTETIAMIEEQNFDIRTITMGISLLDCIDPDINRAAEKIYQKITKAANLVAVGDEIAELGIPVNKRVSVPISLIGAATDATDYVVLAKALDKAAKEIGVDFIGGFSALVQKGYQKGDEILINSIPRALAETDKVCSSVNIGSTKSGINMTAVADMGRRIKETANLSDMGVAKLUVVFANAVEDNPFMAGAFHGVGEADVIINVGSVPGVVKRALEKVRGQSFDDVAETVKKTAFKITRIGQLVQGMASERLGVFEGIVDLSLAPTPAVGDSVARVLEEMGLETVGHTTAALLNDQVKGGVMACNQVGGLSGAFIPVSEDEGMIAAVQNGSLNLKEALMACTAICSVGLDMIAIPEDPTAETIAAMIADAIAIGVINMKTTAVRIIPKGKEGDMIEFGLLLGTAPVMKVNGASSVDFISRGGQIPAPIHSFKNZ

(SEQ. ID. NO. 31)

MTQIIDGKALAALKQQLAEEKTAKLKEETGLVPLGVVILVGDNPASQVYVRNKERSALAAGFRSEVVVPETITQEELLDIAKYNTQPAWHLGVLQPLPKHIDEEAVLLAIDPEKDVGFHPLNMGRWLSGHPVMIPSTPAGIMEMFHEYGIDLEGKNAVIGRSNIVGKPMQALLLAKNATVLTLSRTHNLSKVAAKADILVVAIGRAKFVTADFVKPGAVVIDVGMNRDENGKLCCGDVDYEAVAPLASHITPVPGVGPMTITMLMEQTYQAALRLDRKZ

(SEQ. ID. NO. 32)

MSKFNRIRHLVVLDSVGIGAAPDANNFVNAGVPDGASDTLGHISKTVGLNVPNMAIGLGNIPRETPLKTVAESENPTGYATKLEEV
SLGKDSTMGHWEIMGLNITEPFDFTWNGFPPEILTKIEEFSGRKVIREANKPYSGTAVIYDFGPRQMETGELIYTSADPVLQIAHE
DIIPLDELYRICEYARSITLERPALLGRIIARPYVGEPGNFRTANRRDLAVSPFFPTVLDDKLNEAGIDTYAVGKINDIFNGAGINHDM
GHNKNSNSHGIDLTTKLMGLAEFEKGFSFTNLVDFAFLYDGHRRNAHYGRDCLHEFDERLPEIIAMRENDLLLITADHGNDPTYAG
TDHTREYIPLLAYSAPFKGNGLIPVGHFADISATVADNFGVETAMIGESFLDKLVZ

(SEQ. ID. NO. 33)

MFISAGIVTFLLTVEIPAFIQFYRKAQITGQQMHEDVKHQAKAGTPMGLVFLITSVLVAFFFALFSSQFSNNVMILFILVLYGLVGFLLDFLKVFVRKINEGLNPQOKLALQQLLGGVIFYFLYERGGDILSFGYPVHLGGFYIFFALFWLVGVFSNAVNLTGVDGLASI SVVISLAYSAYVIAVQGQMDILLVILAMIGLGLGGFIFNHPKAVFMGDVGSALGGMLAISMALHQEWTLIIGIVYVFETTSVM MQVSYFKLTGGKRIFRMPVHHHFELGGLSKGNPWSEWKVDFFFFVGVLGALLSLTLAIYLMLZ

(SEQ. ID. NO. 34)

LFKKNKDILNIALPAMGENFLQMLMGMVDSYLVAHGLIAISGVSVAGNIITIYQAIFIALGAAISSVISKSIGQKDQSCLKAYHVTEA
LKJITLLSFSLLGFLSIFAGKEMIGLLGTERDVAEESGGLYLSVGGSVLGLLMSTSGLALIRATHNPRLPLYVSFLSNALNILFSSLAIFV
LDMGIAVGAWGTIVSRVLGVGLWLWSQLKLPYKGPTFGDLKELLTLAPAGERLMMRAGDVIIALVVSGFTEVAGNAIGEVLT
QFNYPMPAFGVATATVMLLARAVGEDDWKRVASLSKQTFWLSFLMLPLSFISIYLVGVPLTHLYTTDSLAVEASVLTFLSLLGTP
MTTGTVIYTAVWQGLGNARLPFYATSIGMWCIRGTYLMGIVLGWGLPGIWAGSLLDNGFRWLFLRYRQYRMSLKGZ

(SEQ. ID. NO. 35)

MQTQEKGHSQAAVGLQHLLAMYSGSILVPIMIATALGYSAEQLTYLISTDIFMCGVATFLQLQLNKYFGIGLPVLGVAFQSVAPI
MIGQSHGSGAMFGALIASGYIVVLVSGIFSKVANLFPSPITVGSVTTIGLTLPVAGINGNMGNVPEPTGQSLLAATVLIILLINIFTKG
FIKSISILIGLVVGTAIAATMGLVDFDSPVAAPLVHVTPLYFGMPTEFESSIVMMCIIATVSMVESTGVYLALSIDTKDPIDSTRLRNG
YRAEGLAVLVLGGGFNTPTYFGTSQNVGLVKLSGKRLPIYYAAGFLVWLNLGPKFGALAQIPISSVLGGMALVMFGFVSIQGMQIL
ARVDFANNEHNFLIAAVSIAAGVGLNNNSNLFSMPTKLFQMFSSNGIVVASLLAIVLNAVLNHKKKZ

(SEQ. ID. NO. 36)

MKDRIKEYLQDKGKVTVNDLAQALGKDSSKDFRELIKTLSLMERKHQIRFEEDGSLTLEIKKKHEITLKGIFAHKNGFGFVSLGE
EDDLFVGKNDVNAYADGDTVEVVIKKVADRNGTAAEAKIIDIHLSLTIVVQGIVLDQEKPKYAGYIRSKNQKISQPIYVKKPAL
KLEGTEVLKFIDKYPSSKKHDFVVASLDLVVGHSTDVGDIVLEVLESMDIVSEFPEVVAEESVPDAPSQKDMERGLDLRDEITF
TIDGADAKDLDDAVHKALKGNLNEFGVHIAVSYYTVEGSDALKENRATSVYVTDRVVPMLPERLSNGICSLNPQVDRLTQS
AIMEIDKHGRVNVYTITQTVKITSFRMTYSVDNDILAGDEEKRKEYHKIVSSIEMAKLHETLENMRVKGALNFDTNEAKILVDK
QGKPVDIVLRQRGIAERMIESFMLMANETVAEHFSKLDPFIYRIHEEPKAEKVQKFIDYASSFGLRIYGTASEISQEALQDIMRAVE
GEPYADVLSSMMLRSQQARYSEHNHGHYGLAADYYTHFTSPIRRYPDLLVHRMIRDYGRSKEIAEHFEQVPIEATQSSNRERRA
IEAEREVEAMKKAEMYMEEVYGEELYDAVVSSIVKFGFLFVELPNTVEGLIHITNLPEFYHFNERDLTRLGEKSGITFRVGQOIRVERA
DKMTGEIDFSVFPSEFDVIEKGLKQSSRSGRGRDSNRRSDKKEDRKSGRSNDKRKHSQKDKKKGKPFYKEVAKKGAKHGKG
RGKGRRTKZ

(SEQ. ID. NO. 37)

MGTTGFTIIDLILIVYLLAVLVAGIYFSKKEMKGKEFFKGDGSPWYVTSVSIFATMLSPISFLGLAGSSYAGSWILWFAQLGMVV
AIPLTIRFILPIFARIIDTAYDYLDRKRFNSKALRIISALLFIYQLGRMSIIMYLPASAGLSVLTGIDINILILMGVVAIVSYTGGKSVL

WTDFIQGVILISGVVLAFLVLIANIKGGFGAVAETLANGKFLAANEKLFDPNLLSNSIFLIVMGSFTILSSYASSQDLVQRFTTQNI
KKLNKMLFTNGVLSLATATVFYLIGTGLYVFYQVQNADSAASNIPQDQJFMYFIAYQLPGITGLILAAIYAQSQTISTGLNSVATS
WTLDIQDVISKNMSDNRTKIAQFSLAVGLFSIGVSIVMAHSDIKSAYEWFNSFMGLVGLLGGVFILGFVSKKANKQGAYAALI
VSTIVMFIFKYLPPATAVSYWAYSISISVSVVSGYIVSVLTGNKVSAPKYTTIHDITEIKADSSWEVRHZ

(SEQ. ID. NO. 38)

MKFSKKYIAAGSAVIVSLSLCAYALNQHRSQENKDNNRVSYVDGSQSSQKSENTPDQVSQKEGIQAEQIVIKITDQGYVTSHGDH
YHYNGKVPYDALFSEELLMKDPNYQLKDADIVNEVKGGYIIVDGKYYVLKDAAHADNRKTDEINRQKQEHVKDNEKVNS
NVAVARSQGRYTTNDGYVFNPADIIEDTGNAIYIPHGGHYHYPKSDLSASELAAKAHLAGKNMQPSQLSYSSSTASDNNTQSV
KGSTSKPANKSENLQSLKELYDSPAQRYSSESDGLVFDPAKIISRTPNGVAIPHGDHYHFIPYSKLSALEEKIARMVPISGTGSTV
NAKPNEVVSSLGSLSSNPLSTSKELSSASDGYIFNPKDIVEETATAYIVRHGDHFHYIPKSQNQIGQPTLPNNSLATPSPLPINPGTS
HEKHEEDGYGFDANRIIAEDESGFVMSHGDHNHYFFKKDLTEEQIKVRKNIZ

(SEQ. ID. NO. 39)

MKKRAIVAVIVLILLGLDQLVKSYIVQQIPLGEVRSWIPNFVSLTYLQNREGAAFSILQDQQLFAVITLVVIGAIWYLNKHMEDSF
WMVGLTLIAGGLNFIDRVSQGFVVMFHLDFNFAINVADSYLTGVILLIAMLKEINGNZ

(SEQ. ID. NO. 40)

MNTNLASFIVGLIIDENDRFYFVQKDGQTYALAKEEGQHTVGDTVKGFAYTDMQKQLRLTTLEVTAQDQFGWGRVTEVRKDLG
VFVDTGLPDKEIVVSLDILPELKELWPKKGDQLYIRLEVDKKDRIWGLAYQEDFQRLARPAYNNMQNQNWPAIVYRLKLSGTFV
YLPENNMLGFHPSERYAEPRLGQVLARVIGFREVDRTLNLSKPRSFEMLENDAMILTYLESNGGFMTLNDKSSPDDIKATFGI
SKGQFKKALGGLMKAGKIKQDQFGTELIZ

(SEQ. ID. NO. 41)

MKDVSFLKKVFKSRLNWIVLALFVSVLGVTFLNSQTANSHSLESRIAANERAINEEKLSQLMSDTSSEEEYQFAKNLD
VQKNLLTRKTEILTLLKEGRWKEAYYLQWQDEEKNYEFVSNPDTASPGLKMGVDRERKIQALYPLNIKAHTLEFPHTGIDQIVWI
LEVIIPLSFVVAIFMLTQLFAERYQNHLDTAHLYPVKVTFIASSLGVGVGVYTVLFICGFSFLVGSISGFGQLDYPYPIYSLVNQ
EVТИКIQDVLFPGLLLAFLAFIVIVEVYLLAIFFKQKMPVFLSLIGIVGLLFGIQTIPQLRQIAHLIPFTYLRSLVEILSGRLPKQIDNV
DLNWSMGMVLLPCLIIFLLGILFIERWGSSQKEFFNRFZ

(SEQ. ID. NO. 42)

MMKFILDIVSTPAILVALIAILGLVLQKKKLDPDIKGGIKTFVGFLVSGGAGIVQNSLNPFQTMFEAHFLSGVVPNNEAIVAVALT
TYGSATAMIMAGMVFNILARFRFYIIFTGHHTLYMACMIAVLVAGFTSLPLILLGGGLALGIMISPAFVQKYMVQLTGNDK
VALGHFSSLGYWLSGFTGSLIGDKSKTEDIKFPKSLAFLRDSTSITLSMAVIYIIVAFAGSEYIEKEISSGTGLVYALQLAQFAA
GVFVILAGVRLILGEIVPAFKGISERLVPSNPKPALDCPIVYTAPNAVLGFISSFVGGLVSMVIMIASGTVVILPGVVPHFFCGATAG
VIGNASGGVRGATIGAFLQGLILISFLPVFLMPVLLGGLFQGSTFSDADFGLSGIILGMLNQFGSQAGIVGLVLIAVMFGVSPFIKKPS
ATEEZ

(SEQ. ID. NO. 43)

MIKTFLSALSVILFSIPIITYSFFPSSNLLNWLSTQPIAQIYAFPLATATMAAILSFLLFFLSFYKKNKQIRFYSGILLLSLILLFGTDK
TLSASNKTTLKLVTWNVANQIEAQHIERIFSHFDADMAIFPELATNIRGEQENQRIKLLFHQVGLSMANYDIFTSPPTNSGIAPVT
VIVKKSYGFYTEAKTFHTRGFTIVLHSRKQNIPDIIALHTAPPLPGLMEIWQKDLNIIHNLASKYPKAIAGDFNATMRHGALAKI
SSHARDNALALPPFERGTWNSQSPKLFNATIDHILLPKNHYYVKDLDIVSFQNSDHRCIFTEITFZ

(SEQ. ID. NO. 44)

MNPQRSWAYVSRKRLRSFILFLILLVLAGISACLTLMKSNKTVESNLYKSLNTSFSIKKIENGOTFKLSDLASVSKIKGLENVSP
ETVAKLKDKDEAVTGEQSVERDDLSAADNNLVSLTALEDSSKDVFTTSSAFNLKEGRHLQKGDSKKILIHEELAKKNGLSLHDKIGL
DAGQSESGKGQTVEFEIIGFSGKKQEKFTGLSSDESENQVFTDYESSQTLLGNEAQVSAARFYVENPKEMDGIMKQVENLALEN
QYQVEKENKAQEIQKDSVATFQTFLTFLYGMILAGAGALILVLSLWLRERYEVGILLALGKGKSSIFLQFCLEVVLVSLGALLP
AFVAGNAITTYLLQTLLASGDQASLQDTLAKASSLSTSILSFAESYVFLVLLSCLSVALCFLFLRKSPKEILSSIZ

(SEQ. ID. NO. 45)

MLHNAFAVYVTRKFFKSIVIFLIILLMASLSLVGLSIKGATAKASQETFKNITNSFSMQINRRVNQGTPRGAGNIKGEDIKKITENKAIE
SYVKRINAIGDLTGYDIELTPTKKNLTADRAKRGFSSLMITGVNDSSKEDKFVSGSYKLVEGEHTNDDKDKILLHKDLAAKG
WKVGDVKVLDSNIYDADNEKGAKETVEVTIKGLDGHNKSATVYSQELYENTAITDIHTAAKLYGYTEDTAIYGDATFFVTADK
NLDDVMKELNGISGINWKSYTLLVKSNSYPALEQSIQMYKMANLLFWGSLFSVLLALLSLWINARRKEVGILLQKQASIL
GQFITESILIAIPALVSAYFLANYTARAIGNTVLANVTSGVAKQASKAAQASNLLGGAEVDGFSKTLSSLDISIQTSDFIIFVLALVLV
VLVMALOSSNLLRKQPKELLLGEZ

(SEQ. ID. NO. 46)

MSQDKQMKAHSVPLLQRVINISSIVGGVGSLFCIWAYQAGILQSKETLSAFIQQAGIWPGLFIFLQILQTVVPIPGALTSVAGVFIYG
HIIGTIINYIGIVIGCAIIFYLVRLYGAAFVQSVVKRTYDKYIDWLDKGNRDRFFIFMMIWPISPADFLCMLAALTKMSFKRYMTII
ILTKPFTLVVYTYGLTYIIDFFWQMLZ

(SEQ. ID. NO. 47)

MRNMWVVIKETYLRHVESWSFFFVISPFLFLGISVGIGHLQGSSMAKNNKVAVVTTVPSVAEGLKNVNGVNFDYKDEASAKEAI
KEEKLKGYLTIDQEDSVLKAVYHGETSELENGIKFETVGTNLNEQNLNRSTASLSQEQEKRLAQTIQFTEKIDEAKENKKFIQTOIAA
GALGFFLYMILITYAGVTAQEVASEKGTKIMEVVFSSIRASHYFYARMMALFLVILTHIGIYVVGGLAAVLLFKDLPFLAQSGILDH
LGDAISLNTLLFILISLFMYYVLA AFLGSMVSRPEDSGKALSPMLIMGGFFGVTALGAAGDNLLKIGSYIPFISTFFMPFRTINDY
AGGAEAWISLAITVIFAVVATGFIGRMYASLVLTQDLDGIWKTFRKRALSYKZ

(SEQ. ID. NO. 48)

MTETIKLMAHTSVRKFKEQEIPQVDLNEILTAQMASSWKNFQSYSVIVVRSQEKKDALLYELVPQEAIRQSAVFLFGDNR
KGARLHTDTFQPQPGVEGLLISSVDAALAGQNALLAELGYGGVIIGLVRYKSEEVAELNLPDITYSVFGMALGVPNQHHDMKP

RLPLENVVFEEYQEQQSTEAIQAYDRVQADYAGARATTSWSQLAEQFGQAEPSSTRKNLEQKKLLZMLKLIAIVGTSKRSTNR
QLLQYMQKHFDTKAELVEIKAIPVFNKPDKQVPAEILEIAAKIEEADGVIIGTPPEYDHSIPA VLSALAWLSYGIYPLNKPIMT
GASYGTLGSSRAQLQLRQILNAPEIKANVLPDEFLLSHSLQAFNPSDLVDLDVIKKLDAIFDDFRIFVKITEKLRNAQELLRKDAED
FDWENLZ

(SEQ. ID. NO. 49)

MNTYQLNNNGEIPVLGFCTFKAKDGEAAYRAVLEALKAGYRHIDTAIIYQNEESVGQAICKDSGVPREEMFVTTKLWNSQQTYEQ
TRQALEKSIEKLGLDYLDLYLIHWPNPKPLRENDAWKTRNAEVWRAMEDLYQEGKIRAIGVSNFLPHHLDALLETTAVPANQV
RLAPGVYQDQVVAYCREKGILLEAWGPFGQGELFDISKVQEIAAHGSVAQIALAWSLAEGFLPLPKSVTSRIQANLDCFGIEL
SHEERETLKTIAVQSGAPRVDDVDFZ

(SEQ. ID. NO. 50)

MRCKMLDPIAIQGLGPLAIRWYALCIVTGLILAVYLTMKEAPRKIIIPDDILDILFILVAFPLAIGARLYYVIFRFDYYSQNLGEIFAIWN
GGLAIYGGGLITGALVLYIFADRKLINTWDFLDIAAPSVMIAQSLGRWGNFFNQEAYGATVDNLDYLPGFIRDQMYIEGSYRQPTFL
YESLWNLLGFALILIFRRKWKSLLRGGHTAFYLIWYGFGRMVIEGMRTDSLMMFFGRVSQWLSVVLIGLGIMIVYQNRKKAPYYIT
EEENZ

(SEQ. ID. NO. 51)

MGKLSSILLGTVSGAALALFLTSKKGKQVCSQAQDFLDDLREDPEYAKEQVCEKLTEVKEQATDFVLKTKEQVESGEITVDSILAQ
TKSYAFQATEASKNQLNNLKEQWQEKAALDDSEEIVIDITEEZ

(SEQ. ID. NO. 52)

MTKKLIFWGSMFLSLSILLTIYAWIFYPMEIQWLNLTNRVYLKPETIQYNFHILMNLYTNPFSQVLQMPDFRSSAAGLHHFAVV
KNLFHVLQVLVALVTPSFYVFVNRIKKDFSLYRKSLLALVLPVMIGLGGVLIGFDQFTLFHQILFVGDDTWLFDPAKDPVIMI
LPETFFLHAFLLFFALYENFFGYLYLKSRRKZ

(SEQ. ID. NO. 53)

MTYHFTEYDIIVIGAGHAGVEASLAASRMGCKVLLATINIEMLAFCMPNCPSIGGSAGIVVREVDALGGEAKTIDKTYIQMKML
NTGKGPAVRALRAQADKELYSKEMRKTVENQENLTLRQTMIDEILVEDGKVVGVRTATHQEYAAKAVVTTGTALRGEIIIDLK
YSSGPNHSLASINLADNLKELGLEIGRFTGTGPPRKASSINYDVTEIQPGDEVPHFSYTSRDEDVYKDQVPCWLTYTNGTSHEIIQ
NNLHRAPMFTGVVKGVGPRYCPSEDKIVRFADKERHQLFLEPEGRNTEEVYQGLSTSLPEDVQRDLVHSIKGLEAEMMRTGY
AIEYDMVLPHQRLATLETKKISGLTAGQTNGTSGYEEAAGQGIIAGINAALKIQGKPELILKRSRGYIGVMIDDVLVTKGTEPYRLL
TSRAEYRLILRHNDADMRLTEMGREIGLVDDERWARFEIKKNQFDNEMKRLDSIKLKPVKETNAKVEEMGFPLTDAVTAKEFLR
RPEVSYQDVVAFIGPAAEDLDDKIELIETEIKYEGYISKAMDQVAKMCRMEEKRIPANIDWDDIDSATEARQKFKLINPETIGQAS
RISGVNPADISILMVLEGKNRSISKTLQSKZ

(SEQ. ID. NO. 54)

MTKVQLLVDDDEEHILKLLDYHLSKEGFSTQLVTNCRKALALAETEPFDIILPQLDGMEVCKRLRAKGVKTPIMMVSAKSD
EDFKVLALELGADDYLTKPFSPRELLARVKAVLRTKGEQEGDDSDNIADDSWLFGTLKVYPERHEVYKANKLLSLTPKEFESDK
NPFPEVFKVSKVTAQZ

(SEQ. ID. NO. 55)

MTTFKDGFWLGGAVAHHQLEGGWQEGGKGIVADVMTAGRHVAREITLGVLEGKYYPNHEADFYHRYKEDIALFAEMGFKC
FRTSIAWTRIFPKGDELEPNEEGLQFYDNLFDECLKNGIEPVITLHFEMPYHLVTEYGGWKNRKLIDFFARFAEVVFKRYKDKVK
YWMFTNEINNQANYQEDFAPFTNSGIVYEEDGNREAIMYQAAHYELVASARAVKIGHEINPDFQIGCMIAMCPIYPVTCNPDKDILM
AMKAMQKRYYFADVHVLGKYPEHIFKYWERKGIVDFTAQDKEDLLGTVDYIGFSYMSFAIDSHRENNPYFDYLETEDLVKN
NYVKASEWEWQIDPEGRLYALNWFTDHYHPLFIVENGFGAIDQVAADGMVHDDYRIEYGAHIREMKA VVEDGVDLMGYTP
WGCIDLVSAGTGEMRKRYGFIYVDKDDNGKGSYRSPKKSFGWYKEVISSNGESVEZ

(SEQ. ID. NO. 56)

MDQQNGLFGLENHVMGPMGKLAQFKVVRITAAGMAAVPFTIVGSMFLVFSILPQAFSFWPIVADIFSASFDFKFTSLYMVANYA
TMGSLSLYFVLSLAYELTKIYAAEELMNPLNGALLALMAFVMTVPQIFDGGMMKTVTSLKEGAVIDGWAMGNVVARFGTT
GIFTAIIMAIVTVLIVYRMCVKHNWVIKMPEAVPEGVSRGFTALVPGFVVAFVVFINGLIVAMGTDIFKVIAPFGFVSNLTNSWIGL
MIYLLTQLLWIVGIGANVFAFSPALIANMAENAAGGHFAVAGEFSNMVFIAGGSGATLGLCLYIAFASKSEQLKAIGRASVVP
ALFNFINEPLGFLPIYNNPALAIPFILAPMVTTATIYYVANSLNFKPIIAQVWPPTVGIGAFLGTADLRAVLVALVCAFAAFLVYLPFIR
VYDQKLVKEEQGIZ

(SEQ. ID. NO. 57)

MKKFYVSPIFPILVGLIAFGVLSTFIIVNNNLLTVLFLVGGYVFLFKKLRYHYTRSDVVEQIQYVNHQAEESLTALLEQMPVGVM
KLNLSSEVEWFNPYAEILITKEDGDFDLEAVQTIKASVGNPSTYAKLGEKRYAVHMDASSGVLYFVDVSREQAITDELVTSRPV
IGIVSVDNYDDLEDETSESIDIQSIVANFISESEKHMMSRRVSMDRFLFTDVTLEGMLNDKFSVIDAFREESKQRQLPLTLS
MGFSYGDGNHDEIGKVALLNLNAEVRRGGDQVVVKENDETKNPVGFGGSAASIKRTRTRTRAMMTAISDKRISVDQVFVVGHK
NLDMDALGSAVMQLFASNVIENSYALYDEEQMSPDIERAVSFILEKEGVTKLLSVKDAMGMVNTRSLLVHDHKTAITLSEFY
DLFTQTIVIDHRRDQDFPDNAVITYIESGASSASELVTTELQFQNSKKNRLSRMAGMMLDTKNFTSRVTSRTFDVASYL
RTRGDSIAIQAIAATDFEEYREVNELIQLQRKLGSDVLLIAEAKDMCYDTVVISKAADAMLAMSGIEASFVLA KNTQGFISIARS
SKLNVQRIMEELGGGGHFNLAAAQIKDVTLSEAGEKLTEIVLNEMKEKEKEEZ

(SEQ. ID. NO. 58)

MKEKNMWKELLNRAGWILVFLLAVLLYQVPLVVTISITLKEVALLQSGLIVAGLSIVLALFIMGARKTKLASFNFSFRAKDLAR
LGLSYLVIVGSNILGSILLQLSNETTTANQSQINDMVQNSSLISSFFLLALLAPICEEILCRGIVPKKIFRGKENLGFVVGTVFALLHQP
SNLPSLLIYGGMSTVLSWTAYTQRLEMSILLHMHVNGIAFCLLALVVIMSRTLGISVZ

(SEQ. ID. NO. 59)

MKEKNMWKELLNRAGWILVFLLAVLLYQVPLVVTSILTLKEVALLQSGLIVAGLSIVVLALFIMGARKTKLASFNFSFRAKDLAR
LGLSYLVIVGSNILGSILLQLSNETTTANOSQINDMVQNSSLISFFLLALLAPICEEILCRGIVPKKIFRGKENLGFVVGTVFALLHQ
SNLPSLIIYGGMSTVLSWTAYKTQRLEMSILLHMIVNGIAFCCLALVVIMSRTLGISVZ

(SEQ. ID. NO. 60)

MDTQKIEAAVKMIIAEGEDANREGLQETPARVARMYQEIFSGLGQTAEEHLSKSFEIIDDNMVVEKDIFFTMCHEHHFLPFYGRA
HIAYIPDGRVAGLSKLARTVEVYSKKPQIQERLNEVADALMDYLGAKGAFVVAEHCMSMRGVKPGTATLTTVARGLFETD
KDLRDQAYRLMGLZMKDLFKLKRQAFRKECLGYLRYVLNDHFVLFLLVLLGFLAYQYSQOLLQHFPEHWPILLFVGITSVLLW
GGTATYMEAPDKLFLVGEEEIKLHLKRQTGISLWFVWLVTFLFLFAPLFLAMGYGLPVFLLYVLLGVGVKYFHFCQKASKFFT
ETGLWDYVISQESKRKQVLLRFFALFTQVKGISNSVKRAYLDFILKAVQKVPKGWQNLYLRSYLRNGDFAISLRLLLSLA
QVFIQAWIATAVVVLFNYLQLFQLLALYHAFDYQYLTQLFLDKGQKEKGLQEVVRGLSFVLLVELVVGILTQEKLALLALLG
AGLVLLVLYLPYQVKRQMHDZ

(SEQ. ID. NO. 61)

MRKSIVLAADNAYLIPPLETTIKSVLYHNRDVDFYILNSDIAPERWFKLLGRKMEVNSTIRSVHIDKELFESYKTGPHINYASYFRFFA
TEVVEDSRVLYLDSIIVTGELATLFIDEILKGYSIGAVDDVYAYEGRKSGFNTGMLLMDVAKWKEHSIVNSLLEAAEQNQVWHL
GDQSILNHYFEDNWLADKTYNMVGIDIYHLAQECERLDDNPPTIVHYASHDKPWNTYSISRLRELWWVYRDLWDSEIAFQRSD
LNYFERSNQSKKQVMLVTWSADIKHLEYLVQRLPDWHFHLAAPCDCSEELTSLSQYTNVTYQVNVLHSRIDWLDDSIVYLDINT
GGEVFNVVTRAQESCKKIFAFDITRKSMDGlyDGIFSVERPDDLVDRMKNEIEZ

(SEQ. ID. NO. 62)

MTKIYSSIAVKGLFTSFLFLFYVLGSRIILPFVVDLNTKDFLGGSTAYLAFSAALTGGNRLSLSIFSVGLSPWMSAMILWQMFSFSKRL
GLTSIEIQDRRKMYLTLIAVIQSLAVSLRPLVQSSYSAILVVLMLNTILLIAGTFFLWVLSLDNLASMIGGSIILLSSMVLNIPQDV
LETFTQVTHIPTGIIVLLALLTUVFSYLLALMYRARYLVPVNKGHLNRFKRSYLEIMLNPAAGGMPYMYVMSFLSPAYLFILLGFIF
PNHSGLAALSKEFMVGKPLWVYVYISVFLFSIIAFVTMNGEIIDARMKKSGEIYGIYPGADTSRFINRLVLRFSVIGGLFNVIMA
GGPMLFVLFDEKLLRLAMIPGLFMFGGMIFTIRDEVKALRNETYRPLIZ

(SEQ. ID. NO. 63)

MSSLSDQELVAKTVEFRQRLSEGESLDDILVEAFAVVREADKRILGMFPYDVQVMGAIVMHYNVAEMNTGEKTLTATMPVYL
NAFSGEVMMVTPNEYLSKRDAEEMGQVYRFLGLTIGVPTEDPKKEMKAEKKLIYASDIYTTNSNLGFYLDNLNASNEEGKF
LRPFNYYIIIDEIDDILLDSAQTPLIAGSPRVSQNSYYAIIDLVTTLVEGEDYIFKEEKEEVWLTGAKSAENFLGIDNLYKEEHASF
ARHLVYAIRAHKLFTKDKDYIRGNEMVLVDKGTGLMEMTKLQQGLHQIAEKEHVKLSPETRAMASITYQLFKMFNKISGMT
GTGKVAEKEFIETYNMSVVRPTNRPQRIDYPDNLITYITLPEKVKYASLEYIKQYHAKGNPLLVFVGSEMSQLYSSLLFREGIAHN
LNANNAAREAQHISESGQMGAVTVATSMAGRGTIDKLGKVVAELGGLIVIGTERMESQRIDLQIRGRSGRQGDGMSKFFVSEDD
VIKFGPSWVHKKYDQVQDMTQPEVLKGRKYRKVLVEKAQHASDAGSARRQTLVEAESMNQIRDIVYKERNRLIDGSRDLE
DVVVDIERYTEEVAADHYASRELLFHFIITVNTISFHKEVPDFYIDVTDKTAVRSMKQVIDKELSEKELLNQHDLYEQFLRLSLLK
AIDNWVEQVDYLQQLSMAIGGQSASQKNPIPEYYQEAYAGFEAMKEQIHADMVRNLLMGLVEVTPKGEIVTHFPZ

(SEQ. ID. NO. 64)

MI GTFAA ALVAV LANFIVPIEITPNSANTEIAPPDGIGQVLSNLLKLVDNPVNALLTANYIRILSWAIFGIAMREASKNSQUELLKT
ADVT SKI VEW IIN LAP F G IL GL VF KT IS DK G VG SL AN Y G ILL V LL V TT M LF V AP V V N P L A F F M R R N P Y PL V N C L R V S G T A F F T R
SSATNIPVN MKL CH D L G L NP DT Y SV SI PL G ST IN MAG V A IT IN LL T LA A V N T L G I P V D F A T A F V L S V V A I S S C D A S G I A G G S L L I P V
ACSLFGISNDIAIQIVGVGFIVGVIQDSCTEALNSSTDVLFTA VAEYAATRKZ

(SEQ. ID. NO. 65)

MSISQRTTKLILATCLACLLAYFLNLSSAVSAGIALLSLSDTRRSTLKLARNRLFSMALLAIGVLAFLSGFHIWSLGLYLAFYVPL
AYKMGWEIGITPSTVLVSHLLVQUESTPDLVNEFLFAIGTGALLVNLYMPSREEEIQHYHTLVEEKLKDILQRFKYYLSRGDGR
NRAQLVAELDTLLKEALRLVYLDHSDHDFHQTDYHIIHYFEMRQRQSRILRNMAQQINTCHLAASESLILAQLFSKIAGQLSQTNPA
SDLLDEIERYLEVFRNRSRSLPKTREEFETRATLLQLLREAKTFIQVKVDFYQKYZQZ

(SEQ. ID. NO. 66)

MEIMSLAIAVFAVIIGLVIGYVSISAKMKSSQEEAELMLLNAEQEATNLRGQAEREADLLVNEAKRESKSLKKEALLEAKEEARKY
REEVDAEFKSERQELKQIESRLTERATSLDRKDDNNTSKEQTLEQKCEQISDRAKNLDAREEQLEEVERQKEAEALERIGALSQAEAR
DIILAQTENEMLTREASIREAEQEVKERSDKMAKDILVQAMQRIAGEYVAESTNSTVHLPPDTTMKGRIIGREGRNIRTFESLTGVD
IIDDTPEVVTLSGFDPRREIARMTMEMLLKDGRHPARIEELVEKRNQREIDNKIREYGEAAAAYEIGAPNLHPDLMKIMGRQLQRTSY
GQNVLRHSIEVAKLAGIMASELGENAALARRAGFLHDIGKAIDHEVEGSHVEIGMELARKYKEPPVVNTIASHHGVDVAESVIAV
IVAAADALSAARP GARSE SLES YIKRLHDLEELIANGFEGVQTSFALQAGREIRIMVNP G KIKDDKVTILAHKVRKKIENNLDYPGNI
KVT VIRELRAVDYAKZ

(SEQ. ID. NO. 67)

MMLKPSIDTLLDKVPSKYSLVILEAKRAHELEAGAPATQGFKSEKSTLRALEEIESGNVTIHPDPEGKREA VRRRIEEKRRKEEEK
KIKEQIAKEKEEDGEKIZ

(SEQ. ID. NO. 68)

MSAYQLPTVWQDEASNQGAFTGLNRP TAGARFEQNLPKGEQAFQQLYSLGTPNGVKV TILLEELLEAGFKEAAYDLYKIAIMDGDQ
FGSDFVKLNPNSKIPALLDQSGTENVRVFESA HILL YLA EKFGAFLPSNPVEKVEVNLWFWQAGA APFLGGGFHFFNYAPEKLE
YPI NRFTMEVKRQLDLDKELAQKPYIAGNDYTIA DIAIWSWYQQLVQGNLYQGSAKFLDASSYQNLVKA EKIANRPAV KRGZ
EV TYTEIKZ

(SEQ. ID. NO. 69)

LASLITSIIMFVYVGFDVLRDTIQKILSREETVIDPLGATLGIISAAIMFVYLYNTRLSKKSNSNALKAAKDNLSDAVTSLGTAIAILA
SSFNYPIVDKLVAIIITFFILKTAYDIFIESSFLSDGFDRLLEDYQKAI M EIPKISKVKSQRGRTYGSNIYLDITLENPDLSVFESHEI
ADQVESM LEE RFGVFD TDVHIEPAPIPEDEILDNVYKKLLMREQLIDQGNQLEELLDDFVYIRQDGEQMDKEAYKT KELNSA IK
DIQITSISQKTKLIC YELDGI IHTS I WRRHETWQNI FHQETKZ

(SEQ. ID. NO. 70)

MTIKLVATDMGTFLDGNGRFDMDRLKSLLVSYKEKGIVFAVASGRGFLSLEKLFAGVRDDIIIFIENGSLVEYQGQDLYEATMS
RDFYLATFEKLKTSPYDINKLLTGGKGSYVLDVDETYLKVSQHNEYIQKVASLEDITDDIFKFTTNFTEETLEDGEAWVNEN
VPGVKAMTTGFESIDIVLDYVDKGVAIVELVKKLGITMDQVMAGCDNLNDLHMMQVVGHPVAPENARPEILEAKTVIGHHKER
SVIAYMEGLZ

(SEQ. ID. NO. 71)

MADIKLIALLDGTLLTDKRLTDRTKETLQAARDRGIKVVLTGRLKAMDFFLHELGTDGQEDEYTITFNGGLVQKNTGEILDK
TVFSYDDVARLYEETEKLSSLPLDAISEGTVYQIQSDQESLYAKFNPALTFVPVFEDLSSQMTYNKCVTAFAQEPLDAAIQKISPELF
DQEYIEFKSREMILLEWSPKNVHKATGLAKLISHLCIDQSQVMACGDEANDLSMIEWAGLVAMQNAVPEVKAANVVTPMTNDE
EAVAWAIEEYVLKENZ

(SEQ. ID. NO. 72)

MESLLILLIANLAGLFLIWQRQDRQEKhLSKSLEDQADHLSQLDYRFQARQASQDQKDLEVVSDRLQEVRIELHQGLTQV
RQEMLDNLNQLTRDKTDQRLQALQESNEQRLEMQRTVEEKLEKTLQASFETVSKQLESVNRGLGEMQTVDGVGALNKV
LSGTKTRGILQEQLQGQIIEDEIMTPAQERYEATVENSERVEYAIKLPGQGDQEYVYLPIDSKFPADYYRLEEAETGDKDEIERC
RKSLLASVKRFARDIRNKYIAPPRTNFVGVLVPTEGLYSEIVRNPVFFDLRREEQIIVAGPSTSALLNSLSVGFKTLNIQKSADHIS
KTLSAVKTEFGKFGGILVKAQKHLQHASGNIDELLNRRTIAERTLRHIELSEGEPALDLLHFQENEEEYEDZ

(SEQ. ID. NO. 73)

MKISHMKKDELFGFYLIKSADLRQTRAGKNYLAFTFQDDSGEIDGKLWDAQPHNIEAFTAGKVVMKGRREVYNNTPVNQIT
LRLPQAGEPNDPADFKVSPDVKEIRDYMSQMIFKIEPNVWQRIVRNLYTKYDKEFYSYPAKTNHHAFETGLAYHTATMVRL
ADAISEVYQLNKSLLYAGIMLHDLAKVIELTGPDTQTEYTVRGNLLGHIALIDSEITKTVMELGIDDTKEEVVLLRHVILSHHGLL
YGSPVRPRIMEAEIIMIDNLDASMMMMSTALALVDKGEMTNKIFAMDNRSFYKPDLZ

(SEQ. ID. NO. 74)

MSEKAKKGFKMPSSYTPLLIAIMAVLTWFIPAGAFIEGIYETQPQNPQGIWVDVLMAPIRALGTHPEEGSLIKETSAAIDVAFFILM
VGGFLGIVNKTGALDVGIASIVKKYKGREKMLLVLMLFALGGTTYGMGEETMAFYPLLPVPMMAVGFDLTLGVIAIILGSQIGC
LASTLNPATGIASATAGVGTGDGIVLRLFWVTLTALSTWFVYRYADKIQKDPTKSLVSTRKEDLKHFNVVESSVESTLSSKQK
SVLFLVLTFLMVLSPFWTDLGVTFDDFTWLTLGPVIGNVGSSTSALGTWYFPEGAMLFAFMGILIGVIYGLKEDKIISFMNG
AADLLSVALIVAIARGIQVIMNDGMITDTILNWGKEGLSGLSSQVFIVVTVYIFYLPMSFLIPSSGLASATMGIMAPLGEFVNVRPSLII
TAYQSASGVNLIAPISTGIVMGALALGRINIGTWWKFMGKLVVAAIVVTIALLLGTFLPFLZ

(SEQ. ID. NO. 75)

MSNSFVKLLVSQLFANLADIFFRVTIIANIYIISKSVIATSLVPILIGISSFVASLLVPLVTKRLALNRVLSLSQFGKTILLAILVGMFTV
MQSVAPLVTLFVVAISILDGFAAPVSYAIVPRYATDLGKANSALSMTGEAVQLIGWGLGGLLFATIGLLPTTCINLVYIISFLML
FLPNAEVEVLESETNLEILLKGWKLVARNPRLRFLFSANLLEIFSNTIWVSSIIIVFVTELLNKTESYWGNSNTAYSIGIIISGLIAFRLS
EKFLAAKWEPLQLFTPNLKTIQNPCLSLDPGWFLSPNGCFLDKKEFPLYGISVEKNTKRKETHMNSLSPNHHFQNKSFYQLSFDDG
HLTQYGGLIFQELFSQLKLKERSKYLVNTDQRRCYRSQDSDSLVLQFQLTGYTDYACKELSADAYFPKLEGGQLASQPTLS
RFLSLRTDEETVHSLRCLNLELVEFFLQFHQLNQLIVDIDSTHFTTYGKQEGVAYNAHYRAHGYHPLYAFEGKTYCFNAQLRPGN
RYCSEEADSFTPVLERFNQLLFRMDSGPATPKLYDIEKTGQYYLIKLLKNTVSLRGDLSLPCPQDEDLTILPHSAYSETLYQAGS
WSHKRVCQFSERKEGNLFYDVISLVTNMSTSGTSQDFQLYRGRGQAENFIKEMKEGFFGDKTDSTSLIKNEVRMMMSCIAYNLY
LFLKHLAGGDFQTLTIKRFRLFLHVVGKVRTGRKQLLKLSSLYAYSELFALSIRKVNLPVYEPRRKASLMMHZ

(SEQ. ID. NO. 76)

MMEFFQQQLPHLEPYGNPQYFVYVIAATLPFIGLFFKKRFAWYEVLVSLFFIVTMLVGGKTNQALALGIYLCWEIILLLFYKHYRK
KDGGKWFVYLVFSFLSLPIIFVVKVQPAINGTQSLLGFLGISYLTFRSVGIVIELRDGVIKDFTLWEFLRFLFMPFTSSGPIDRFKRFNEN
YQAIPERDELMMDMDESVRYIMWGFYKFILAHVLGETLLPPLKNLALQSGGFLNLYALA VMYTGLELFFDFAGYSMFALAIISNL
MGIRSPINFNKPFLSRDLKEFWRNWRHMSLFWFRDFVFMRMVMVLTRKKVFKNRNTSSMAYIVNMLIMGFWHGVWTYYIAG
LFHGLGLVINDAWVRKKKTLNKERKAGKAALPENRWIQLLGMVVTFHVVMLSFLIFSGFLNNLWFKZ

(SEQ. ID. NO. 77)

MLKRLWMIFCPVLIAGLLVFLIFFYPTEMHHNLGAEKRSAVAATTIDSFKERSQVKRALSDPNVRFPFFGSSEWLRFDGAHPAVL
AEKYNRSYRPYLLGQGGAASLNLQYFGMQQMLPQLENKQVYVVISPCWFSKNGYDPAAFQQYFNGDQLTSFLKHQSGDQASQYA
ATRLLQQFPNVAMKDLVQKLASKELSTADNEMIELLARFNERQASFFGQFSVRYVNYDKHVAKYLKILPDQFSYQAIEDVVKA
DAEKNTSNEMGMENYFYNEQIKKDLKLLKDSQKSFTYLKSPBEYNDLQLVLTQFSKSKVNPIFIIPPVNKKWMNYAGLREDMYQ
QTVQKIRYQLESQGFTNIADFSKDGGEPEFFMKDTIHLGWLWAFDKAVDPFLSNPTPAPTYHLNERFFSKDWATYDGDVKEFQZ

(SEQ. ID. NO. 78)

MEKNLKALKQTTDQEGPAIEPEKAEDTKTVQNGYFEDAAVKDRTLSDYAGNWQSVPFLEDTFDQVFDFYKAKLTGKMTQA
KAYYTKGYHTDVTKINITDNTMEFVQGGQSKKTYKYVGKILTYKKGNRGVRFLFEATDADAGQFKYVQFSDHNVAPVKA
FHIFGGTSQEALFEEMDNWPTYYPDNLSGQEIAQEMLAHZ

(SEQ. ID. NO. 79)

MKDGHLLAHHIRLLNCRFKLQLSQDPEALYRGEQGKILA VLWNSETGCATATDIALATGLANNTLTTMIKKLEEQKLVIVSPCGK
DKRKKYLVLTELGKSQKEVGHRSQKLDTIFYKGFSEEEIHQFEGFQERILANLKEKGNEVZ

(SEQ. ID. NO. 80)

MTNLIATFQDRFDWLTALSQHQLQLSLLTLLAILLAIPAVFLRYHEKLADWVLQIAGIFQTIPSALLGLFIPLMGIGTLPA
LYAIFPILQNTITGLKGIDPNLQEAGIAFGMTRWERLKKFEIPLAMPVIMSGIRTAAVLIGTATLAALIGAGGLGSFILLGIDRN
LIGALSSAVLIAIFNFLKVMEMAKLRTIFSGFALVALLGLSYSPALLVQKEKENLVIAGKIGPEPEILANMYKLLIEENTSMTATV
KPNFGKTSFLYEALKKGDDIDIYPEFTGTVTESLLQPSPKVSHPEQVYQVARDGIAKQDHLAYLKPMPSYQNTYAVAVPKKIAQEY
LKTISDLKKVEGQLKAGFTLEFDREDGNKGLQSMYGLNLNVATIEPALRYQAIQSGDIQITDAYSTDAELERYDLQVLEDDKQLF
PPYQGAPLMKEALLKKHPELERVLNTLAGKITESQMSQLNYQVGVEGKSAKQVAKEFLQEGLKKZ

(SEQ. ID. NO. 81)

MMHTYLQKKIENIKTTLGEMSGGYRRMVAAMADLGSGTMKAIWDDLFAHRSFAQWIYLLVLGSFPLWLELVYEHRIVDWIGMI
CSLTGIIICVIFVSEGRASNYLFGLINSVIYLILALQKGFYGEVLTLYFTVMQPIGLLWVWIYQAQFKKEKQEFVARKLDGKGWTKYL
SISVLWWLAFGFIYQSIGANRPYRDSITDATNGVGQILMTAVYREQWIFWAATNVFSIYLWWGESLQIQGKYLIYLINSLVGWYQ
WSKAQKNTDNNZ

(SEQ. ID. NO. 82)

MRNMKAKYAVVVAFFLNLTIAIVEFIAGGVFGSSAVLADSVHDLGDAIAIGISAFLETISNREEDNQYTLGYKRFSLGALVTAVI
LVTGSVVLVILENVTKILHPQPVNDEGILWLGIIAITINLLASLVGKGKTKNESILSHFLEDTLGWVAVILMAIVRFTDWYILDPLL
SLVISFFILSKALPRFWSTLKIFLDAVPEGLDIKVQKSGLERLDNVASLNQLNLWTMDALEKNAIVHVCLKEMEHMETCKESIRIFL
KDCGFQNITIEIDADLETHQTHKRKVCDLERSYEHQHZ

(SEQ. ID. NO. 83)

MIEYKNVALRYTEKDVLRDVNLIQEDGEFMVLVPGSGSGKTTMLKMINRLLEPTDGNIYMDGKRIKDYDERELRLSTGYVLQAIA
LFPNLTVANIALIPEMKGSKEEIKKTEELLAKVGLPVAEYGHRLPSELGGEQQRVGIVRAMIGQPKIFLMDEPFSALDAISRK
QLQVLTKELHKEFGMTTIFVTHDTDEALKLADRIAVLQDGERQVANPETILKAPATDFVADLFGGSVHDZ

(SEQ. ID. NO. 84)

MSAVAISAMTKVMQETHGNPSSIHGHRQAGKLLREARQELAQLLRTKPQHIFTSGGTEGNNTIIGYCLRHQEQGKHIITTAIEH
HAVLETIDYLVQHFGFEATIIQPNQEITAQQIQKALRDDTILVSTMVFVNNETGNLLPIAEIGQILKQHPAAHVDAVQAIGKIPIHSE
ELGIDFLTASAHHKFHGPKGICFLYASSMDFDSYLHGGDQEKKRAGTENLPAIVGMVAALKEDLEKQEEHFQHVQNLETAFLAEL
EGIQYYLNRGKHHLPYVLNIGFPQGKNDLLLRLLAGISISTGSACTAGVVQSSHVLEAMYGANSERLKESSLRISLSPQNTVEDLQ
TLAKTLKEIIGGZ

(SEQ. ID. NO. 85)

MLFKLSKEIQLGSLRSPARRIFLSFALVILLGSLLLSLPFVQVESSRATYFDHFTAUSAFCVTGLSTLPVAHTYNIWGQIICLLIQ
IGGLGLMTFIGVFYIQSKQKLSLRSRATIQDSFSYGETRSLRKFVYSIFLTTFLVESLGAILLSFRLIPQLGWGRGLFSSIFLAISAFCA
GFDNLGMLTSFLAQFTDLLVNLVIAGLITGGLGFMVWFDFLAGHVGRKKKGRHLHFTKLVLLLTIGLLFGTATTLFLEWNNAQTIG
NLPVADKVLVSFFQTVTMRTAGFSTDITYTQAHPVTLLIYILOMFLGGAPGGTAGGLKITTFVLLV FARSELLGLPHANVARTIAP
RTVQKSFSVFIIFLMSFLIGLILLGITAKGNPPFIHLVFETISALSTVGTANLTPDLGLKLAALSIVIMPLMFMRIGPLTFVSLADYHPE
KKDMIHYMKA DISIGZ

(SEQ. ID. NO. 86)

MSDRTIGILGLGIFGSSVLAALAKQDMNIIADDHAERINQFEPVLARGVIGDITDEELLRSAGIDTCVTVVVATGENLESSVLA
CKSLGVPTVIAKVKSQTAKVKLEKIGADSVISPEYEMGQSLAQTLFHNSVDVFQLDKNVSVEMKIPQSAGQSLSKLDLRGKYN
LNILGFREQENSPLDVEFGPDDLLKADTYILA VINVQYLDTLVALNSZ

(SEQ. ID. NO. 87)

MKLLSIAISSYNAAYLHYCVESLVICGEQVGILIINDGSQDQTQEIAECLASKYPNIVRAIYQENKCHGGAVNRLGVEASGRYFKV
VDSDDWVDPRAYLKILETLQELESKGQEVDFVTNFVYEKEGQSRKMSYDSVLPVRQIFGWDQVGNFSKGQYTMMHSIYRT
DLLRASQFZ

(SEQ. ID. NO. 88)

MKFNPNQRYTRWSIRRLSVGVASVVASGFFVLVGQPSSVRADGLNPTPGQVLPEETSGTKEGDLEKPGDTVLTQAKPEGVTGN
TNSLPTTERTEVSEETSPSSLDTLFKEDEAQKNEELTDVLKETVDTADVDGTQASPAETTPEQVKGGVKENTKDSIDVPAAYLEK
AEGKGPFAGVNQVIPYELFAGDGMLTRLLKASDNAPWSDNGTAKNPALPPEGLTKGKYFYEVDLNGNTVGKQGQALIDQLR
ANGTQTYKATVKVYGNKDKADLTLNIVATKNVDININGLVAKETVQKA VADNVKDSIDVPAAYLEKAGEGPFATGVNHV
ELFAGDGMLTRLLKASDKAPWSDNGDAKNPALSPLEGENVTKGQYFYQVALDGNVAGKEKQALIDQFRANGTQTYSATVNVY
GNKDGKPDLDNIVATKVTININGLISKEVQKA VADNVKDSIDVPAAYLEKAGEGPFATGVNHVIPYELFAGDGMLTRLLKA
SDKAPWSDNGDAKNPALSPLEGENVTKGQYFYQLADGNVAGKEKQALIDQFRANGTQTYSATVNVYGNKDGKPDLDNIVATK
KVTININGLISKEVQKA VADNVKTVSMFQQPTZ

(SEQ. ID. NO. 89)

MKLKSYILVGYIISTLLTILVVFWAVQKMLIAKGEIYFLLGMTIVASLVGAGISLFLLPVFTSLGKLKEHAKRVAAKDFPSNLEVQG
PVEFQQLGQTNFEMSHDLQVFSDFSLEESEREKGLMIAQLSHDIKTPTSIQATVEGILDGIIKESEQAHYLATIGRTERLNKLVEELN
FLTLNTARNQVETTSKDSIFLDKLLKIECMSEFQFLIEQERRDVHLQVIPESARIEGDYAKLSRILVNLVDNAFKYSAPGKTLLEVVAKL
EKDQLSISVTDEGQGIAPEDLENIFKRLYRVETSRNMKTGGHGLGLAIARELAHQLGGEITVSSQYGLGSTFTLVNLNGSENKAZ

(SEQ. ID. NO. 90)

MFGQTAQHGLTNSLKDFWIFLLNIGPQLAFFCQMLRCSRSEVEQGTGNHRREFNMIQQIFSHFGMTHLGQIKLVYQESIDLELLVNA
LNHHLLIDRLVLTNPQNTIEIDRQIVHGLDLLKGRKDKEIIDIKSMFRQLELASTQQICPINQRVHHGILAFGEISDLVPAKNLPNRQDZ

(SEQ. ID. NO. 91)

MEHLATYFSTYGGAFFAALGIVLAVGLSGMGSAYVGKAGQSAALLKEQPEKFASALILQLLPGTQGLYGFVIGILIWLQLTP
PLEKGVAYFFVALPIAIVGYFSAKHQGNVAVAGMQLAKRPKEFMKGAILAAMVETYAILAFVVSFTLKV

(SEQ. ID. NO. 92)

MLKSEKQSRYQMLNEELSFLLEGETNVLANLSNASALIKSRFPNTVFAGFYLFDGKELVLGPFQGGVSCIRIALGKGVCGEAHFQ
ETVIVGDVTTLYNYISCDSLAKSEIVVPMKNGQLLGVLDLDSSEIEDYDAMDRDYLEQFVAILLEKTAWDFTMFEKSZ

(SEQ. ID. NO. 93)

MSVLEIKDLHVEIEGKEILGVNLTLKTGEAAIMGPNGTGKSTLSAAIMGNPNEYVTGEVLFDGVNILELEVDERARMGLFLAM QYPSEIPGITNAEFLRAAMNAGKEDDEKISVREFITKLDEKMELLNMKEEMAERYLNEGSGGEKRNEILQLMLEPTFALLDEID SGGLDIDALKVVSKGVNAMRGEGFGAMIITHYQRLNYITPDVVHVMMEGRVVLSSGPELAARLEREGYAKLAEELGYDYKEELZ

(SEQ. ID. NO. 94)

MPYKRQRFSFMSALKLDSLYMAVVADHSKNPHHQGKLEDAEQISLNNTCDGVINLSVKFDAEDRLEDIAFLNSGCTISTASASM MTDAVLGKTKQEILELATIFSEMVGQKDERQDQLGDAFLSGVAKFPRIKCATLAWNALKTIENQEKQZ

(SEQ. ID. NO. 95)

MKIQDLLRKDVMLLDQATEKTAVIDEMIKNLTDHGYVTDFTFKEGILAREALTSTGLGDGIAMPHSKNAAVKEATVLFAKSNK GVDYESLDGQATDLCMIAPEGANDTHLAALAEQSQYLMKDGFADKLQRQATSADQVIELFDQASEKTEELVQAPANDSGDFIVA VTACTTGTIAHTYMAQEALQVAAEMVGIVKVTENGASGVGNQLTAEDIRKAKAIIAADKADEMDFDGKPLINRPVADGIRKTE ELINLALSGDETYVRAANGAKAATASNEQSLGGALYKHLMSGVSQMLPVIGGGIMIALAFIDGALGVPNENLGNLGSYHELA SMFMKIGGAAGFLMLPVAGYVAYSIAEKPLVAGFVAGAIKEGFAFGKIPYAGGEATSTLAGVSSGFLGALVGGFIAGALVL AIKKYVKVPRSLEGAKSILLPLLGILTGFVMLAVNIPMAINTAMNDFLGGGGSAVLLGIVLGGMMAVDMGGPVNKAYV FGTGTLAATVSSGGSVAMAAMAGGMVPPPLAIFVATLLFKDKFTKEERNGLTNIMGLSFITEGAIPFGAADPARAIPSFILGSAVA GGLVGLTGIKLMAPHGGIFVIALTSNALLYLVSVLVGAIVSGVYGYLRKPQAZ

(SEQ. ID. NO. 96)

MANKNTSTTRRPSKAELERKEAIQRMLISLGLIAILLIFAAFKLGAAGITLYNLIRLLVGSLAYLAIFGLLIYLFFFKWRKQEGLLSGF FTIFAGLLLIFEYLWVKYGLDKSVLKGTMAQVVTDLTGFRTTSFAGGGIIGVALYIPTAFLFSNIGTYFIGSILLVGSLLVSPWSVY DIAEFFSRGFAKWEGHERRKEERFVKQEEKARQKAECLEARQEETEACALLDLPVDMETGEILTEEAVQNLPPIPEEKWVPEII LPQAELKPKPEQEDDSDEDVQVDFSKEALEYKLPLQLFAPDKPKDQSKEKKIVRENKILEATFASFGIKVTVERAEIGPSVTKEYE VKPAVGVRVNRSNLSDDLALALAACKDVRIEAPIPGKSCLIEVPNSDIATVSRFLWEQSQTKAENFLEIPLGKAVNGTARAFDSLK MPHLLVAGSTGSGKSVAVNGIIASILMKARPQVKFMMVDPKMVELSVYNDIPHLLIPVVTNPRKASKALQKVVDEMENRYELFA KVGVNRNIAFGNAKVEEFNSQSEYKQPLPFIVVIVDELADLMMVASKEVEDAIRLGQKARAAGIHILATQRPSVDVISGLIKANV PSRVAFAVSSGTSRSTILDENGAEKLLGRGDMFLFKPIDENHPVRLQGSFISDDDVERIVNFIKTQADADYDESFDPGEVSENEGEFS DDGAGGDPFLFEAKSLVIETQKASASMIQRRLSVGFNRATLMELEIAGVIGPAEGTKPRKVQQZ

(SEQ. ID. NO. 97)

MSYFKKYKFDKSQFKLGMRTFKTGINFLVLLIFGFFGWKGLQIGALTAVFSLRESFDESVFHGTTSRILGNSIGGLYALVFLNNTFF HEAFWVTLVVVPICTMLTIMTNVAMNNKAGVIGVAAMLIITLSPGETILYVFWVLETFMGVFVAIIVNYDIRLIRLFLEKKEKZ

(SEQ. ID. NO. 98)

MNKSEHRHQLIRALITKNKIHTQAEQALLAENDIQVTQATLSRDIKNMNLSKVREEDSAYYVLNNGSISKWEKRLELYMEDALV WMRPVQHQVLLKTPGLAQSFSGSIIDTLSFPDAITLCGNDVCLIICEDADTAQKCFEELKKFAPPFFEEZ

(SEQ. ID. NO. 99)

MKSINKLNALSYMGIRVNLNIIPILTGTYYVARVLDRTDYGYNFNSVDTILSFLPFATYGVYNYGLRAISNVKDNKKDLNRTFSSLFYLC IACTILTTAVYILAYPLFFTNDNPVKKVYLVGMQIQLAQIFSIEWVNEALENYSFLYKTAFRILMLVSIFLFLVKNEDIVVYTLVMSL STLINYLISYFWIKRDIKLVKIHLSDFKPLFLPLTAMLVTFANANMLFTFLDRLFLVKTGIDVNVSYYTIAQRIVTVAGVVTGAIGVSV PRLSYYLGKGDKEAYVSLVNRGSRIFNFFIIPLSFGMLVLPNAIILYGESEYIGGGILTSLFATRITIILALDTILGSQILFTNGYEKRIT VYTVFAGLLNLGLNSLLFNHIVAPEYLYLTTMSETSLVFYIIFIHRKQLIHLGHIFSYTVRYSLFSLSFVAIYFLNFVYPDMVIN LPFLINTGLIVLLSAYISLLVFTKDSIFYEFLNHVLAALKNKFKKSZ

(SEQ. ID. NO. 100)

MELFMKITNYEYKLKKSGLTNQQILKVLEYGENVDQELLLGDIADISGCRNPAVFMERYFQIDDAHLSKEFQKFPSFSILDDCYPW DLSEIYDAPVLLFYKGNLDDLKFPKVAVVGSRACKSGAKSVEKVIQGLENELVIVSGLAKGIDTAAHMAALQNGGKTAIVGVTG DVFVYKANKRQLDYIGNDHLVLSEYGPGEQPLKFHFPARNRIIAGLCRGVIVAEAKMRSGSLITCERAMEEGRDVFAPGSILDGLS DGCHHLIQEGAKLVTSGQDVLAEEFZ

(SEQ. ID. NO. 101)

MQLTQEDAKQIELEILDYIDTLCKKHNNINYIINYGTIGAVRHEGFIPWDDDDIDLSMPREDYQRFINIFQKEKSKYKLLSLETDKNY FNNFIKITDSTTKIIDTRNTKTYESGIFIDIFPIDRFDDPKVIDTCYKLESFKLSSFSKHKNIVYKDSLLKDWRITAFLWLLRPSPRYFA NKEIKEIQKYSRENGQYMAFIPSKFKEKEVFPSTGTFDKTIDLPFENLSPAPEKFDTILTQFYGDYMTLPPEEKRFYSHFHAYKLED Z

(SEQ. ID. NO. 102)

MIKINHLLTITQNKDLRDLVSDLTMITQDGEKVIIIGEEGNGKSTLLKILMGEALSDFTIKGNIQSDYQSLAYIPQKVPEDLKKKTLHD YFLFLSDIDLDSYISLYRLAELHFDSNRFASDQEQIGNLSGGEALKIQLIHELAKPFEIFLDEPSNLDLETVDWLKGQIQKTRQTVIFIS HDEDLFLSETADTIVVHRLVCHRKEAETLVEHLDYDSYSEQRKANFAKQSQQAANNQRAYDKTMKRRVKQNVENTRALRATKDS TAGRLLAKKMKTCLSQEKRYEKAQSMSQTQKPLEEIQQLFFSDIQPLPASKVLUQLEKENLSIDDRVLVQKLQLTVRGQEKGIIIGP NGVGKSTLLAKLQRLLNDKREISLGFMPCDQYHKKLQLDLSPIAQLSKTGEKEELQKIQSHLASLNFSPYEMQHQIERSLSGGQQGKL LLLDLVLRKPNFLLDPEPTRNFSPTSQPQIRKLFATYPGLITVSHDRRLKEVCSIYRMTTEHGLKLVNLEDZ

(SEQ. ID. NO. 103)

MKPKTFYNLLAEQNLPLSDQQKEQFERYFELLVEWNEKINLTAITDKEEVYLKHFYDSIAPILQGLIPNETIKLLDIGAGAGFPSLPM KILYPELDVTIIDSINKRINFLQLAQEELDNGVHFYHGRAEDAFAQDKNFRFAQYDFVATARAVRMQVLSELTIPYLKVGKLLALK ASNAPEELLEAKNALNLFSKVEDNLSYALPNRDPRYITVVEKKKETPNKYPRKAGMPNKRPLZ

(SEQ. ID. NO. 104)

MSIKLIAVDIDGTLVNSQKEITPEVFSAIQDAKEAGVKVVIATGRPIAGVAKLDDQLRDEGVDYVVTFNGALVQETATGHEIIESL TYEDYLDMEFLSRKLGVHMHAITKDIYTANRNIGKYTVHESTLVSMPIFYRTPEEMAGKEIVKCMFIDEPEIILAAIEKIPAEFYER

YSINKSAPFYLELLKKNVDKSAITHLAEKLGLTKDETMAIGDEENDRAMLEVGNPVMENGNEIKKIAKYITKTNDSEGVAH
AIRTWVLZ

(SEQ. ID. NO. 105)

MTWIILGVIALIVIFIVSYNGLVKNRMQTKEAWSQIDVQLKRRNDLLPNLIETVKGYAKYEGSTLEKVAELRNQVAAATSPAEM
KASDALTRQVSGIFAVAESYPDLKASANFVKLQEELNTENKISYSRQLYNSSNVNKLETFPSNIIAGMFGKAADFLQTPEEE
KSVPKVDFSLGDZ

(SEQ. ID. NO. 106)

MLFDQIASNKRKTWILLLVFFLLALVGAVGYLFIIRSGLGGLVIAJIIGFIYALSMIFQSTEIVMSMNGAREVDEQTAPDLYHVVE
DMALVAQIPMPRVFIIDDPALNAFATGSNPQNAAVAATSGLLAIMNREELEAVMGHEVSHIRNYDIRISTIAVALASAITMLSSMAG
RMWWGGAGRRRSDDDRDGNGLEIIMLVVSLLAIVLAPLAATLVQLAISQRREFLADASSVELTRNPQGMINALDKLDNSKPM
RHVDDASSALYINDPKKGFFQKLFYTHPPISERIERLKQMZ

(SEQ. ID. NO. 107)

MKLN1QEIRKQSEGLENFEQTLDLVDDLRARNQEILDVKDILA VGKVQYEDRMYFLDYQLSYTIVLASSRSMEPVELVESYPVTEVF
MEGATNQLDQEVLDDLVLPIENGELDAESVSDNILLNIPIKVLTAEEEAGQGFISGNDWQIMTEEYQAQKAVKKEENSPFAGL
QGLFDGDEZ

(SEQ. ID. NO. 108)

MKRLALVVFSGGQDSTCLFWVMQHYETVEAVTFAYGQRHHLEIQITREIAKEQGIRHHILDMSLLGQITAQPDFATIHISYIPDK
LCVESKSLKLYLFSYRNHGDFHENCINTIGKDLVNLLDPRYLEVWGKFTPRGGISIDPYNYGKQGTKEYEGLAEQRLFQHDLYPEK
IDNRZ

(SEQ. ID. NO. 109)

MTETVEDKVSHSITGLDILKGIVAGAVISGTVATQTKVFTNESAVLEKTVKTDALATNDTVVLGTISTSNSASSTSLSASESASTS
ASESASTSASTSASTSASESASTSISASSTVGSQTAATEATAKKVEEDRKKPASDVASVTNVNLQSYAKRRKRSVDSIEQ
LLASIKNAAVFSGNTIVNGAPAINSLNIAKSETKVTGEGVDSVYRPIYYKLKVTNDGSKLTFTVTVYVNPCTNDLGNISSMRP
GYSYNSGTSTQMTLTLGKPGVKNYITDKNGRQVLSYNTSTMTCQSGYTWNGNAQMNGFFAKKGYGLTSSWTVPITGT
DTSTFTPYAARTDRIGINYFNGGGVKESSSTSQSLSQSKLSVSASQASASASTSASASASTSASASASTSASASASTSASVAST
SASASASTSASASASTSASESASTSASASASTSASASASTSASASASTSASESASTSASASASTSASASASTSASASASTSA
SGSASTSTSASASTSASASASASASESASTSASASASTSASESASTSASASASTSASASASTSASASASTSASASASTSASAS
TSASESASTSASASASTSASASASTSASASASTSASVSASTSASASASTSASASASTSASESASTSASASASTSASASASTS
ASASASTSASESASTSASASASTSASESASTSASASASTSASASASTSASASASTSASASASASASESASTSASE
SASTSTSASASTSASESASTSASASASTSASASARQVRPQVHNRHQVRQPQQVVLHQLQHQRVRHLQHQVPRVPL
QRQPVRQLQQVPLQSQHQVLPQHQVPLQQVHQVPLQQVHQVPLQQVHQVPLQQVHQVPLQQVHQVPLQQVHQVPLQQVHQVPL
QPVHQSQSQVPRQLPHQQVPRQLQQVHQVPLQQVHQVPLQQVHQVPLQQVHQVPLQQVHQVPLQQVHQVPLQQVHQVPL
RLNRHQVRPLQQVHQVPLQQVHQVPLQQVHQVPLQQVHQVPLQQVHQVPLQQVHQVPLQQVHQVPLQQVHQVPLQQVHQVPL
QHQQAPQLQQVPRQPQQVPRQLQQVPRQPQQVPRQPQQVPRQPQQVPRQPQQVPRQPQQVPRQPQQVPRQPQQVPRQPQQVPRQP
HQSQQVPRQFRINKCLGFSKYZ

(SEQ. ID. NO. 110)

MGVETWFYSSICWLAIGLGSVWKFPYMTAANGGGGFLILFLISTILIGFPLLAEFALGRSAGVSAIKTFGKLGKNNKYNFIGWIGA
FALFILLSFYSSVIGGWILVYLGIEFGKLFQLGHTGDYAQLFTSIISNPALGAQAAFILLNIFIVSRGVQKGIERASKVMMPLLIVFV
FIIGRSLSLPNAMEGVYFLKPDFS KLTS TGLLY ALGQSF FALS LGVT VMLTY ASYLDKKTNLVQSGSISIVAMNISIMAGLAIFQZ
SPNFIQSEGGPSLLFIVLPQLFDKMPFGTIFYVLFLLFLFATVTFSVMLIEINVNDNITNDNSKRAKWSVILGIFTVFGIPSALSYGV
MADVHIFGKTFDAMDFLVSNLLMPFGALYLSFTGYIFKKALAMEELHLDERAWKQGLFQVWLFLRFVSSFQSSLWSSLPNL
CNQKGLEZ

(SEQ. ID. NO. 111)

MLKKWQLKDVLILLAFLSIFFGGVFVGSGYVYNILSLLTPLGLQAFANEILFGLWCMAAPIAAIFVPRVGSATIGEVLAALAEVLYG
SQFGLGALLSGFVQGLSEFGIVTKNRYESWLSLTANSIGITLVSFVYEYIKLGYYAFSLPFVLSLLVVFISVYFFCTILVRAIVKL
YHQFATGGKAZ

(SEQ. ID. NO. 112)

MVKVATQTPIISLFLILSLETSFIPSIALTLSVVAFCILFMLYYRRFKMLAWMIILAILPSFANYWA VQLHGDA SQA VMLGTRAFVT
VCIGLVFVSSVSLKELLLYLAQKGLRSWSYALIVVNFSPLIQKQEI KSLKEACLLRGQELHFWSPLIYSKVLMTVFRWRHLYLRAL
SAHGYDEHAQLKNSYRTFYIPKKTKLIYLLFLQLTSLFLZ

(SEQ. ID. NO. 113)

MRKHQLQVHKLTILSMMIALDVVLTPIFRIEGMAPMSSVNVILAGIMMGPVYALAMATVTAFIRMTTQGIPPLALTGATFGALLAG
LFYKYGRKFHYSLAGEILGTGIGSIVSYPVMVLFTGSAAKLSWFIYTPRFFGATLIGTAISFIAFRFLIKQEFFKKVQGYFFSERIDZ

(SEQ. ID. NO. 114)

MQEFTNPPIGSSSLIHCITNEISCEMLANGILALGCKPV MADD SREVLDFTKQSQALFINLGHLSAEKEKAIRMAAS YANQSSLPMV
VDAVGVTSSIRKSLVKDLDYRPTVLKGNMSEIRS LVGLKHGVGV DASA KQDQETEDLLQVLKDWCQTPGMSFLVTGP KDLV
VSKNQVAVLNGC TELDWITGTGDLVGA LTA VFLSQGK TGFEAS CLA VSYLNIAAEKIVVQGMGLEEF RYQVLNQLSLLRDEN
WLDTIKGEVYEZ

(SEQ. ID. NO. 115)

MNHKIAILSDVHG NATALE AVIADAKNQGASEWLLGDIPLPGP GANDLV ALLKDLPTASVRGNWDDR VLEALDGQYGL ED PQ
EVQLLRMTQYLMERMDPATIVWLRSPLLEKKEIDGLRFSISHNLPDKNYGGD L VENDTEKFDQLDAETDV AVYGHVHKQ LL
RYGSQGQQIINPGSIGMPYFNWEALKNHR SQYAVIEVEDGEL NIQFRKVAYDYEAELELA KSKG L PFIEM YEELRRDD NYQGHNL
ELLASLIEKHGYVEDVKNFFDFLZ

(SEQ. ID. NO. 116)

MNVNQIVRIIPTLKANNRKLNETFYIETLGMKALLEESAFLSLGDQTGLEKLVLEEAPSMRTRKVEGRKKLARLJVVKENPLEIEGILSKTDSIHRLYKGQNGYAFEIFSPEDDLILIAEDDIASLVEVGEKPEFQTDLASLSKFEISMEHLHPTDIESFLESSEIGASLDIFPAQGQDLTVNDNTWDLMSMKFLVNELDIASLRQKFESTEYFIPKSEKFLGKDRNNVELWFEEVZ

(SEQ. ID. NO. 117)

MKWTKIICKIEEQIEAGIYPGASFAYFKDNQWTFYLGQSDPEHGLQTEAGLVYDLASVSKVVGVGTCTFLWEIGQLIDIRLVIDFLPESDYPDTIRQLLTHATDLPFPINRNDLLTAPELKEAMFHNLRRSQPAFLYSVHFLLGFLERIFNQDLDVILKDQVWKPGMTETKFGPVLA VPTVRGVEAGIVHDPKARLLGRHAGSAGLFSTIKDLQIFLEHYLADDFARDLNQNFSPLDDKERSLAWNLEGDWLDHTGYTGTFIMWNRQKQEATIFLSNRTYEKDERAQWILDRNQVMNLIRKEEZ

(SEQ. ID. NO. 118)

MMKKTYNHILVWGVIFYSICIVCFCTPQEQTGVGVTGPGIQHLGRVLFLTPNSLWKLGEVDIGQLCWIFLQNILNVFLFFPLIFQLLYLFNLRKTKVLLFSFLVSLGIECTQLI LDFFFDFNRVFEIDLWTNTLGGYLAWLLYKRLHKNKVRNZ

(SEQ. ID. NO. 119)

MKIPLLTFARHKFVYVLLTLLFLALVYRDVLMTYFFFIDIHAPDLAKFDGQAIKNDLLKSALDFRILQFNLCGFYQSFIIPIIIVLLCGFOYIELKNCVNLRLSISGREVSYQGLKRKLTQVASIPCLILYVTVLIIAITYFFGTFSPGLGWSLSDGSGLQRLLDGEIKSYLFFTCVLLIGIFINAIFYLQIVDVGNVNTRSAITYLMLFWLGSMLLYSALPYYMPVMTSLMQASYGDVSLMKLFTPYIYIVPYMVLEKYEDNVZ

(SEQ. ID. NO. 120)

MFKVLCVKVKGAFMLPIALPAAGLLLIGGGALSNPPTIATYPILDNSIFQSIFQVMSSAGEVVFSNLSLLCVGLCIGLAKRDKGTAALAGVTGVLVMTATIKLVLFMAEGSAIDTGVIGALVVGIVAVYLNHNRYYNIQLPSALGFFGGSRVPIVTSFSSILIGFVFFVIWPFFQQLVSTGGYIQCAGPIGTFYGLMRLRSGAVGLHHIYPMFWYTELGGVETVAGQTVVGAQKIFFAQIADLAHSGLFTEGTRFFA GRFSTMMFGLPAACLAMYHSVPKRNKKYAGLFFGVALTSFITGITEPIEFMFLFVSPVLYVVHAFLDGVSSFIADVLNISIGNFTSG GVIDFTLFGIQLQNAKTNWVLQIPGLIWSVLYIIFRWIFQFNVLTGPRGEEVDSKEISESADSTSNTADYLKQDSLQIIRALGGSN NIEDVDACVTRLRVAVKEVNQVDKALLQIGAVDVLEVKGQIYGAKAILYKNSINEILGVDDZ

(SEQ. ID. NO. 121)

MKFRKLACTVLAGAAVLGLAACGNSSGSKDAAKSGGDGAKTEITWWAFPVFTQEKTGDGVGTYEKSIIEAFEKANPDIKVKLETIDFKSGPEKITTAIEAGTAPDVLFDAPGRIIQYKGNGKLAELNDLFTDEFVKDVNNENIVQASKAGDKAYMYPISSAPFYMANKK MLEDAGVANLVKEGWTTDFEKFVLKALKDKGYTPGSLFSSGQGGDQGTRAFISNLYSGSVTDEKVKSYTTDDPKFVKGLEKATS WIKDNLINNNGSQFDGGADIQNFANGQTSYTLWAPAQNGIQAQKLEASKVEVVEVPFPSDEGKPALEYLVNGFAVFNNKDDKKVA ASKKFIQFIADDKEWGPKDVRVTGAFPVRTSGFKLYEDKRMETISGWTQYYSPYNTIDGFAEMRTLWFPMQLQSVNGDEKPADA LKAFTEKANETIKKAMKQZ

(SEQ. ID. NO. 122)

MQSTEKKPLTAFTVISTIILLLTBLFIPFYWILTGAFKSQPTDIVERPPQWFPKMPTMENFQQLMVQNPALQWMWNSVFISLVTMFLVCATSSLAGYVLAALKRFYQQRILFAIFIAAMALPKQVVLVPLVRIVNFMGIHDTLWAVIDPLIGWPFVGFLMKQFSENIPTELLESA KIDCGEIRTFWSVAFPIVKPGFAALAIFTINTWNDYFMQLVMLTSRNNLTISLGVATMQAEMATNYGLIMAGAALAAVPIVTVF LVFQKSFTQGITMGAVKGZ

(SEQ. ID. NO. 123)

MKIMFKNFNNILLNRKIVLLLRLVLMAMILINHLLSTAVQKDAVIFFKRELISIFSNDYSEANLEIPKLLNLSLFMVGWLSVILLESDLADHYHHLIRYQSSFFDYTRKRLVVISKFFTQDLFWFLGLPLGIHFKTVALFLLAQLMMLYLLSYLIALISAGAGFSFFLYLFALVGQEWMMDHVTVYLVLLSLLVMLIVSRLEEKFKKGZ

(SEQ. ID. NO. 124)

MGKGEMGKGIVIGLEFDSEVLVNKAPTLQLANGKTATFLTQYDSKTLFAVDKEDIGQEIIGIAKGSIESMHNLPVNLAGARVPGVNGSKAAVHEVPEFTGGVNGTEPAVHEIAEYKGSDSLVTLLKDYTYKAPLAQQALPETGNKESDLLASLGLTAFFLGLFTLGKKREQZ

(SEQ. ID. NO. 125)

MKTFPLLVLGLFCLLPLSVFAIDFKINSYQGDLYIHADNTAEFRQKIVYQFEEDFKGQIVGLGRACKMPSGFIDPHPKIQAACKNG AELADVTSEVTEEADGYTVRVYNPGQEGDIVEVDLVWNLKNNFLYDDIAELNWQPLTDSSIESIEKFHVRGDKGAEKLFHTG KLFRECTIEKSNLQDYTIRLDNLPAKRGVELHAYWPRTDASFARDQGLKGNRLEEFNKIEDSIVREKDQSKQLVTWVLPSILSILLLS VCFYFIYRKRTTSPVKYAKNHRLYEPPMELEPMVLEAVYSTSLEEVSPLVKGAGKFTDQLIQATLLVIDRGNVSISEGDAVGL RLVKEDGLSSFEKDCLNLAFSGKKEETLSNLFADYKVSDSLYRRAKSDEKRIQARGLQLKSSFEELNQMCEGVRKRSFWGLPDYRPLTGGKECALQVGMGALTIPFLFIGFLFLYSLDVHGFLYLPILGFLGLVLSVFFYWLRLDNRDGVNEAGAEVYLLWT SPENMLREIARLDQAELESIVVWNRLLVYATLFGYADKVSHLMKVHQIVENPDINLYVAYGWHSTFYHSTAQMHSYASVANTA STYSSSGSGSSGGFSGGGGGSIGAFZ

(SEQ. ID. NO. 126)

MKKVRKIFQKAVAGLCCISQLTAFSSIVALAETPSPAIGKVVVKETGEGGALLGDAVFELKNNTDGTVSQRTEAQTGEAIFSNIK PGTYTLEAQPPVGYPKSTKQWTVEVEKNGRTTVQGEQVNEREEALSDQYPQTGTYPDVQTPYQIIKVDGSEKNGQHKALNPNPY ERVPEGTLSKRYQVNLLDDNQYGIELTVSGKTVYEQKDKSVPLDVILLDNNSNSMSIRNKNARRAERAGEATRSLIDKITSDE NRVALVTVASTIFDGTETFTVEKGVADKNGKRLNDSLFWNYDQTSFTNTKDYSYKLTKNDKNDIVELKNKVPTEAEDHDGNRLM YQFGATFTQKALMKADEILTQQARQNSQKVIFHITDGVPTMSYPINFNHTFAPSQYQNQLNAFFSKSPNKGDGILLSDFITQATSGEHTIVRGDGQSYQMFTDKTVYEKGAPAAFPVKEKYSEMKAAGYAVIGDPINGGYIWLNWRESILAYPNSNTAKITNHGDPTRWYY NGNIAPDGYDFTVIGIGINGDPGTDEATATSFMQSISSKPNETYNTVTDTTKILEQLNRYFHTIVTEKSIENGTTIDPMGELIDLQLGTDGRFDPADYTLTANDGSRLENGQAVGGPQNDGGGLKNAKVLYDTTEKIRVTGLYLGTDKETVLTLYNVRLNDEFVSNKFYDTNG RTTLHPKEVEQNTVRDFIPKIRDVRKYPEITISKEKLGDFIEFKVNKNDKPLRGAVFSLQKQHPDYPDIYGAIDQNGTYQNVRT GEDGKLTFKNLSDGKYRLFENSEPAGYKPVQNKPIVAFQIVNGEVRDVTSIVPQDIPAGYEFTNDKHYITNEPIPPKREYPRGIGMLPFYLGCMMSGVLLYTRKHPZ

(SEQ. ID. NO. 127)

MKSINKFLTMLAALLTASSLFAATVFAAGTTTSVTVKLLATDGMDMKIANELETGNYAGNKVGVL PANAKEIAGVMFVWT
NTNNEIIDENGQTLGVNIDPQTFLSGAMPATAMKLTEAEGAKFNTANLPAAKYKIYEIHSLSYVGEDGATLTGSKAVPIEIELP
LNDVVDAHYPKNTAEAKPKIDKDFKGKANPDTPRVDKDTPTVNHQVGDVVEYEIVTKIPALANYATANWSDRMTEGLAFNKGTV
KVTVDDVALEAGDYALTEVATGFDLKTDAGLAKVNDQNAEKTVKITYSATLNDAIVEVPESNDVTNFNYGNPNDHGNTPKPNK
PNENGDLTTLKTWVADTAPIPAGEATEFLVNAQTGKVQVTLTTDKNTVTVNGLDKNTYEKFVVERSIKGYSADYQEITTA
IAVKNWKDENPKPLDPTEPKVVTYGKKFVKVNDKDNRLAGAEFVIANADNACQYLARKADKVSQEEKQLVTTKDALARVA
AYNALTAQQQTQKEKEVKDQAAYNAAVIAANNAFEWVADKDNEVVKLVSDAQGRFEITGLLAGTYYLEETKQAGYALL
TSRKFEVTATSATGQGIEYTAGSGKDDATKVVNNKKTIPQTGGIGTIIFAVAGAAIMGIAVYAYVKNKDEDQLAZ

(SEQ. ID. NO. 128)

MTMQKMOKMISRIFFVMAFCSLVWGAHAQAEQEDHTLVLQLENYQEVVSQLPSRDGHRLQVWKLDDSYSYDDRVQIVRDLHS
WDENKLSSFKKTSFEMTFLENQIEVSHIPNGLYYVRSIIQTDASVSYPAEFLFEMTDQTVPLVIVAKKDTMTTKVLIKVDQDHNR
LEGVGFKLVSVARDVSEKEVPLIGEYRYSSSGQVGRFTYTDKNGEIVFTNLPLGNYRFKEVEPLAGYAVTTLTDVQLVDHQLVTI
TVNVQKLPGRNVDFMKVDRNTSLOGAMFKVMKEESGHYTPVLQNGKEVVVTSGKDGRFRVEGLEYGTYLWELQAPTYGV
QLTSPVSFTICKDTRKELTVVVKNNKRPRIDVPTGEETLVYLDACCFFVVWZ

(SEQ. ID. NO. 129)

MSHIYLSIFTSLLLMLGLVNVAQADEYLIGMEAAAPFNWTQDDDSNGAVKIDGTNQYANGYDVQIAKKIAKDLGKEPLVVKT
KWEGLVPALTSGKIDMIIAGMSPTAERKQEIAFSSYYTSEPVLLVKDSAYASAKSLDDFNGAKITSQQGVLYNLIAQIPGAKKE
TAMGDFAQMRQALEAGVIDAYVSEPEALTAEEANSKFKMIQVEPGFKTGEEDTAIAIGLRKNDNRISQINASSETISKDDQVALM
DRMIKEQPAEATTETSSSSQVAKILSENWQQLRRAGITLISIVGTLIIGLALGIVFRTAPLENKVNIVGLQKLVGVWLNVYI
EIFRGTPMIVQSMVIYYGTAQAFGINLDRTLAAIFVISNTGAYMTEIVRGILAVDKGQFEAATALGMTHNQTMRKIVLPQVVRNI
LPATGNEFVINIKDTSVLNVISVVELYFSGNTVATQTYQYFQFTIIAVYFVLTFTVTRILRFIERMDMDTYTTGANQMATEDLKZ

(SEQ. ID. NO. 130)

MTQAILEIKHLKKSYGQNEVLKDISLTVHKGEVISIIGSSGSGKSTFLRSINLLETPTDGQILYHGQNLEKGYDLTQYREKLMVFQ
SFNLFENLNVLENTIVAQTTVLKRERTEAEKIAKENLEKVGMDERYWQAKPKQLSGGQKQRVAIARALSMNPDAILFDEPTSALD
PEMVGEVLKIMQDLAQEGLTMIVTHEMFARDVSHRVIFMDKGVIAEEGKPEDLFTNPKEDRTKEFLQRYLKZ

(SEQ. ID. NO. 131)

MKKYQLLFKISAFTSYLFFFVFSQLSQLTLIVQNYWQFSSQIGNLFWIQNILSLLFIGVMIVVLVKTGHGYLFRIPRKKWLWYSILTVL
LVFQISFNVQTAKHVQSTAEGWAFLIGYSGTNFAELGIYIALFFLVPMLMEELIYRGLLQHAFFKHSRFLDLLLPSILFALPHFSSLPS
LLDIFVFATVGIIFAGLTRYTKSIYPSAVHVINNIVATFPFLLTFLHRVLGZ

(SEQ. ID. NO. 132)

MNKKQWLGLVAVAAGVLAACGNRSSRNAASSSDVKTKAIAVTDGGVDDKSFNQSAWEGLQAWGKEHNLSKDNFTYFQS
TSEADYANNLQQAAAGSYNLIFGVGFALNNAVKDAAKEHTDLNYVLIDDVKDQKNAVSFTADNESGYLAGVAAAKTTTKQV
GFVGGIESVISRFEAGFKAGVASVDPISKVQVDYAGSGFDAAKGKTIAAAQYAAGADIVYQVAGGTGAGVFAEAKSLNESRPEN
EKVVIGVDRDQEAEKGYTSKDGKESNFVLVSTLKQVGTVDISNKAERGEFPGGQVIVYSLDKGVDAVTNLSEEGKKAVE
DAKAKILDGSVKVPEKZ

(SEQ. ID. NO. 133)

MSKKLQQISVPLISVFLGILLGAIVMWIFGYDAIWGYEELFYTAFGSLRGIGEIFRAMGPLVLIGLGFVASRAGFFNVGLPGQALAG
WILSGWFALSHPDMPRPLMILATIVIALIAGGIVGAIPGILRAYLGTSEVIVTImmNYIVLVGNAFIHAFPKDFMQSTDSTIRVGAN
ATYQTPWLAELTGNSRMNIGIFFAIIAVAVIWFMLKTTLGFEIRAVGLNPHASEYAGISAKRTIILSMIISGALAAGLGA
QNVYVQGSSLAIGFNGMAVSLLAANSPIGILFAAFLFGVLQVGAPGMNAAQVSELVSIVTASIIFVSVHYLIERFVKPKQVKGG
KZ

(SEQ. ID. NO. 134)

MGVKKKLKLTLLGLSLLIMTACATNGVTSITAESADFWSKLVYFFAEIIRFLSDISIGVGIILFTVIRTVLLPVFQVQMVASRKM
QEAPRIKALREQYPGRDMESRTKLEQEMRKVFEMGVRSQDSLWPILQMPVILALFQALSRVDFLKTGHFLWINLGSVDTTLVL
PILAAVFTFLSTWLSNKALSERNGATTAMMYGIPVLIFAVYAPGGVALYWTVSNAYQLTYFLNNPFKIIAEREAVVQAQKDL
ENRKRKAKKAAQTKZ

(SEQ. ID. NO. 135)

MVIDPFAINELDYYLVSHFHSDHIDPYAAAILNNPKLEHVFKFIGPYHCGRIWEGWGPKERIIVVKPGDTIELKDMKIHAVESFDR
TCVLTPVNGADETGGELAGLAUTDEEMAQKAVNYIFETPGGTIYHGADSHFSNYFAKHGKDFKIDVALNNYGENPVGIQDKMT
SIDLLRMAENLRTKVIIPVHYDIWSNFMASTNEILEWLWKMRKDRLQYDFHIFIWEVGGKYTPQDQHLVEYHHPRGFDDCFEQDS
NIQFKALLZ

(SEQ. ID. NO. 136)

MFLSGWLSSFANTYIHDLGVLFDPFLNAFESAIAAPLVEEPLKLLSLVFVLALIPVRKLKSLFLGIASGLGFQMIKDIGYIRTDLP
EGFDFTISRLERIISGIAASHWTFSGLAVVGVYLLRAYKGQKVGKKQGLIFLGLALGTHFLNSPFVELETPLAIPVVTIALYGF
YHAYCFVEKHNELMTZ

(SEQ. ID. NO. 137)

MKVEPRCDVLSRMSHFFIRILIMELQELVERSWARQAYHELEVKHHD SKWTVEEDLLALSNDIGNFQLVMTKQGRYYDETPYT
LEQKLSENIWWLLELSQRLLDIDILTEMENFLSDKEKQLNVRTWKZ

(SEQ. ID. NO. 138)

MLDWKQFFLAYLRSRSRSLFIYLLSLAFLVLLFQFLFASLGIYFLYFFFLLCCFVTILFFTWDILVETQVYRQELLYGEREAKSPLEIALA
EKEAREMELYQQRSKAERKLTDLDDYYTLWVHQIKTPIAASQLVAEVVDRQLKQQLEQEIFKIDSYTNLVLQYLRLESFHDDLV

LKQVQIEDLVKEIIRKYALFFIQKGLNVNLHDLDEIVTDKWLLVVIEQIISNSLKYTKEGGLEIYMDQELCIKDTGIGIKNSDVL
RVFERGFSGYNGRLTQQSSGLGLYLSKKISEELGHQIRIESEVGKGTTRVIQFAQVNVLVEZ

(SEQ. ID. NO. 139)

MELNTHNAEILLSAANKSHYPQDELPEIALAGRSNVGKSSFINTMLNRKNLARTSGKPGKTQNNFFNIDDKMRFVDVPGYGYAR
VSKKEREKGCMIEEYLTTRENLRRAVSVLVDLRHDPSADDVQMYEFLKYYEIPVIIVATKADKIPRGKWNKhesAIKKLNFDPS
DDFILFSSVSKAGMDEAWDAILEKLZ

(SEQ. ID. NO. 140)

MTTKQLHLVITGMMSGAGKTVAIQSFEIDLGYFTIDNMPPALLPKFLQLVEIKEDNPKLA
VVDMRSSRSFFFSEIQA
VLDELENQDGL
DFKILFLDAADKELVARYKETRRSHPLAADGRILDGIKLERELLAPLK
MSFGFKYGPIDADLVFDVRFLPNYYLPELRNQTGVDEPVYDYVMNHPESEDFYQHLLALIEPILPSYQKEGKS
HRSVAFAKRLAQDLSKNWSVNEGHRDKDRRKETVNRZ

(SEQ. ID. NO. 141)

MRKPKITVIGGGTGSPVILKSLREKDVEIAAVTVADDGGSSGELRKNMQQLTPPGDLRNVLVAMSDMPKFYEKVFQYRFS
AFAGHPLGNLIIAGLSEMCGSTYNAMQLLSKFFHTGKIYPSSDHPLTLHA
ASRRVVQTI
VPQEYMSNSRFDEYLVQEHDFVGLCKQVS
IQVKKZ

(SEQ. ID. NO. 142)

MKNLIKLLIIRLIVNLADSVFYIVALWHVSNYSSSMFLGIFIAVNYLPDLLL
WVMSLVFISVMASSISYVIEDV
ANIE
QIVGVFLFLNGSSWLVAIV
AISIPYFLFSYIFYTDNGLKEFSIYZ

(SEQ. ID. NO. 143)

MMSNKNK
SEWIGAVGYLGGGLAFSIYVKEKENIY
V
KZ

(SEQ. ID. NO. 144)

LKKPIIEFKNVSKV
LFPHMNV
MQYELRELQQLGITFV
AVDGGM
TEEEFDARIEEY
V
KZ

(SEQ. ID. NO. 145)

MKSMRILFL
V
VPNSVK
V
KZ

(SEQ. ID. NO. 146)

MVKKIIGMV
NP
KZ

(SEQ. ID. NO. 147)

MKKQAYV
VV
VY
KZ

(SEQ. ID. NO. 148)

MIVSISQGF
LV
AL
A
LIL
KZ

(SEQ. ID. NO. 149)

MKMKV
VIA
KZ

(SEQ. ID. NO. 150)

MIKFDNI
VFDNIA
KZ

(SEQ. ID. NO. 151)

MRHKLNLKDWLIRLGLIWFLVTFIYPNFDLVNVFVKGGESLDAVRVLKSQRALQSIMNSFKLAFLSLIITNVVGILCVLFTEYF
DIKGAKILKLGYMTSLIYGGVVLATGYKFVYGPYGLITKFLQNVIPSLDPNWFIGYGAVLFIMTFSGTANHTLFLNTIRSDYHTIE
AARNMGAKPFTVFRKVVLPTLIPTLFAUTIMVFLSGLSAVAAPMIVGGKEFQTINPMIITFAGMGNRSRDLAALLAIILGIATTILLTIM
NKEKGNNYISISKTКАPLKKQKIASKPWNIIAHIVAYGLFTVFMLPLIFIVLYSFTDPVAIQTGNLTSNFTLENYRLFFSNSAAFSPF
LVSFIYSIIAATTATILAVVVARVVRKHKSRFDFLFEGALLPWLLPSTLLAVSLLFTFNQPQFVLNQILVGSLVILLIAYIVVKIPFSY
RMVRAILFSVDEMEDAARSMGASFYTMMKVIIIPFILPVVLSVIALNFNSLLTDFDLSVFLYHPLAQPLGITIRSAGDETATSNAQA
LVFVYTIVLMIISGTVLYFTQRPGRKVRKZ

Table 3**ID201 - 4106.4 (SEQ. ID. NO. 168)**

ATGATAAAAATCCTAAATTATTAACCAAGTCTTTTAAAGAAGTGGTCAATTCTAGGTGGTGTGGTCTAGTCATTCAATAGCTATTATTTGACCTTCCCTTTTATTATATTCAACTGGAGGGGGAAAAGTTAATGAGAGCGCAAGAGTGTGTTACCGAGTATTAAAGACTAACATCTGATGAAAGATATTGAGATAAGCGGCTCCCTTGTGCATGACTGGATATTAAAGATGAAAGCTATCAAATTATATCGTGTGTTAGATATGTCTGTTAGTACAGCAGATGGTAAACAGGTAAACCGTCAATTGTTACGGGGTGGATGTCTACAAAGAAGCAAAGAATTTTGCTTGTGTTACATCCCATATACATTGGTTACAATTGCTTTCCCTTGTGTTCTTATTTTAACTAAACGC TTGCTCAATCCTTGTGTTACATTCAGAAGTGACTIONTAAAGTCAAGAATTGGTAAACCTGACTGAGTGGACAAATATTGCTTGTGAGAAAGTAG GAAAGATGAGTGGTGAAGTGGAAAACAGATAATGGTATGTGAGCACTGGTGAAGGTTATTATGAGTTGGAAAGTC GTAATGAGCAAATTGAAATTGCAAATCAAAGGTTCTTGTCCGGAGCATCACATGAGTTGAAACCCCTTGTGCC AGTCTTGAATTATCCTAGAGAATATGCAGCATAATATTGGAGATTACAAGATCATCCAAATATATTGCAAAGAGTATAAA TAAGATTGACAGATGAGCCATTAGAAGAAGTACTGGAGTCTCTAAATTCAAGAGTGGACAGAGTGTGAGACCT TGACTGTTAAGCCAGTTAGTAGATATTATCACGTTATCAAGAATTAGTCATCAATAGGTGTTACAATTGAAAATCAA TTGACAGATGCTTACCGGGTGTGATGAGTCTTGGGCAATTGATAAGGTTTGACAAACCTGATTAGTAATGCAATTAAATA TTCAGATAAAATGGGCGTGAATCATATCGAGCAAGATGGCTATCTCTATCAAATCGTACAGATAAGGATGAAAGTCCGGTTGGTCTTAC ATTGTGAATAATATTAGAAAGCTATCAAATGGATTATAGTTCTCCCTATGAACACGGTATGAAATTAGATTAGCTT GTAG

(SEQ. ID. NO. 152)

MIKNPKLLTKSFLRSFAILGGVGLVIHIAIYLTFPFYYIQLEGEKFNESARVFTEYLKTKTSDEIPSLLQSYSKSLTISAHLK RDIVDKRLPLVHLDLIDKGKLSNYIVMLDMVSSTADGKQVTQFVHGVDVYKEAKNILLLYLPYTFLTIASFVFSYFYTKR LLNPLFYISEVTSKMQDDNNIRFDESKRDEVGEVGKQINGMYEHLKVIELESRNQIVKLQNOKVSFVRGASHELKTPLA SLRIILENMQHNIGDVKDHPKYIAKSINKDQMSHLLEEVLESSKFQEWECRETLTVKPVLVDILSRYQELAHSIGVTIENQ LTDATRVVMSRALDKVLTNLISNAIKYSDKNGRVIISEQDGYLSIKNTCAPLSDQELEHLFDIFYHSQIVTDKDESSGLGLY IVNNILESYQMDYSFLPYEHGMEFKISLZ

ID202 - 4106.9 (SEQ. ID. NO. 169)

ATGGATAAAAATTATTAACATATCAGAAAGCGGAGCCTTCGTGCTTGTCTGATAGCACTGAAACCGTCCGCACACTGCTCAAGAAAAACATCAAACCCAAGCTAGCTCAACTGTAGCGCTTGGTCGAACCTTATCGCTAGCCAGATTCTCGCAGCCAATG AAAAAGGAAATACCAAACCTACAGTTAAGGTGTTGGATCTAGCTCTAGGTGCTATTATCACCGTCGCTGATACCAAGGGG AACGTCAAAGGCTATGTTCAAACCTCTGGTGTGACATCAAACAAAGACTCGCAGTGGTAAGTCCTAGTGGACACCTTTGTTGG AAAAGCTCAATTCTCTGTGTTACAGACTCGGAAACTTCAACTTACAGACTCGGAAAGGATCTGCTCATCTCTGGAGAAATCG GTGAAGACCTGCTTTTACCTTACTGAAAGCAACAAACGCCCTCAGGGTCGGCCTCAATGCTCTTTGGACGAGGAAGAC AAGGTCAAGGGTGCAGGGTTCTAGTTCAAGTCTGCCAGGAGCCAAGAAAGAGAGATTGCTCGCTTGAAAACGCAT CCAAGAAATGCCAGTACTCTACTCTCTGAAAGCAGCACCATATCGAAGGCCCTCCTCAAGGCTATCACGGGACGAAG CCTACAAGCGCTTTCTGAAAGAAGAAATCGGTTCCAATGTAAGTGTACTGAGCCATGAACGCTTATGAACGCTTGTGCAATTCTGCCAAACTACTTACAACCT TGATGAAAAGGACCTGGAGGAACTCATTGTCACAAATTCTAA

(SEQ. ID. NO. 153)

MDKIIKTISESAGRAFVLDSTETVRTAQEKHQTQASSTVALGRTLIASQILAANEKGNTKLTVKVLGSSSLGAIITVADTKG NVKGVVNPGVDIKKTATGEVLVGFVGNQFLVITDYGTGNPYNSTPLISGEIGEDLAFYLTESQQTPSAVGLNVLLDEED KVKGAGFLVQVLPAGKKEEIRFEKRIOEMPAISTLLESDHIEALLKAIYGDEAYKRLSEEIRFQCDCSHERFMNALASL PSSDLQEMKEEDHGAETCQFCQTTYNFDEKDLEELIRDKSZ

ID203 - 4115 (SEQ. ID. NO. 170)

ATGAAATCAATAACTAAAAGATTAAGCAACTTTGCAGGAGTAGCTGCCTTGTGAGTATTGCTCCATCTTGTATC TGCTCAAGAAATCATCAACTTACACTGTTAAAGAAGGTGATACACTTTAGAAATCGCTGAAACTCACAACACAACAGTTGAAA AATTGGCAGAAAACAACCCACATGTAACATTCTGATTATGTTGATCAAGAGTTGGTATCGATGGCCTGTAGCGCCT GTTGCAACACCAGCGCCAGCTACTTATGCGGCCAGCGCCTCAAGATGAAACTGTTAGCTCAGCTCAGTAGCAGAAACTCCAGT AGTAAGTGAACACAGTTCAACTGTAAGCGGATCTGAAAGCAGAACGGGAGAAATCACTTGTCAATTCTGCCAAACTACTTACAATTGTA GTATACAGCTACAAATGGACGTTATCGGACGTTACCAATTAA

(SEQ. ID. NO. 154)

MKSITKKIKATLAGVAALFAVFAPSFSVQAESSTYTVKEGDTLSEIAETHNTTVEKLAENNHDNIHLIYVDQELVIDGPVAP VATPAPATYAAPAAQDETUVSAPVAETPVSETVSVTSGSEAEKEWIAQKESGGSIQLQMDVISDVTNZ

ID204 - 4117.1 (SEQ. ID. NO. 171)

ATGAATTAGGAGAATTGGTACAATAAAATAAGAACAGAGGAAGAGTTAATGAAGAAAAGTAAGATTATTTT AGCTCTGCTATTCTTAGCTAGTCAGAGGGTGCAATGGCTAGTGATGGTACTGGCAAGGAAAACAGTATCTGAAAGAAG ATGGCAGTCAGCAGCAAATGAGTGGGTTTTGATACTCATATCAATCTGGTTCTATATAAAAGCAGATGCTAACTATGCT GAAAATGAATGGCTAAAGCAAGGTGACGACTATTCTGACATCTGGTGGCTATATGCCAAATCAGAATGGTAGAAGA CAAGGGAGCCTTTATTATCTGACCAAGGAGATGAAAAGAAATGCTTGGTAGGAACCTCCTATGGTGTGCAACAG GTGCCAAAGTAAGAAGACTGGGCTATGATTCTCAATACAGATGCTTGGTTTATCAAAAGCAGATGGACAGCACGAGAAAG AAAGAATGGCTCAAATTAAAGGAGGAGACTATTATCTGACATACAGCAAGGTTGGCTTTGACAAACAATACCAATCTGGTTTACATCAAAGAAA ATGAAACTATGCTGATAAGAATGGATTTCGAGAATGGTCACTATTATCTAAATCCGGTGGCTACATGGCAGCCAAT GAATGGATTGGGATAAGGAATCTGGTTTATCTCAAATTGATGGGAAAATGGCTGAAAAGAATGGGTCTACGATTCTCA TAGTCAGCTTGGTACTACTTCAAATCCGGTGGTTACATGACAGCCAATGAATGGATTGGGATAAGGAATCTGGTTTATC

TCAAATCTGATGGGAAAATAGCTAAAAAGAATGGGTCTACGATTCTCATAGTCAGCTGGTACTACTCAAATCCGGTGGT
TACATGACAGCCAATGAATGGATTGGATAAGGAATCTGGTTTACCTCAAATCTGATGGGAAAATAGCTGAAAAGAATG
GGTCTACGATTCTCATAGTCAGCTGGTACTACTCAAATCTGGTGGTACATGGCAGAAAATGAGACAGTAGATGGTTATC
AGCTTGGAAAGCGATGGTAAATGGCTGGAGGAAAACCTACAAATGAAATGCTGCTTACTATCAAGTAGTGCTGTTACAGCC
AATGTTATGATTGAGCTGGTAAAGCTTCCATATCGCAAGGAGTGTCTGCTGAGATAAGGATAGAAAAGTGA
TGACAAGCGCTGGTACTTACTATTCTGGTTGTCAGGCTATATGAAAACAGAAGATTTACAAGCGTAGATGCTAGTAAGG
ACTTTATCCCTTATTATGAGAGTGTGGCCACCGTTTATCACTATGTCAGGCTCAGAATGCTAGTATCCCAGTAGCTTCTCAT
CTTCTGATATGGAAGTAGGCAAGAAAATTATCGGCAGATGGCCTGCATTTGATGGTTAAGCTTGAGAATCCCTTCCT
TTCAAGGATTTAACAGAGGCTACAAACTACAGTGCTGAAGAATTGGATAAGGTTAGTTAGGCTAAACATTAACAATAGCC
TTTGGAGAACAGGGCTACTTTAAGGAAGCCGAAGAACATTACCATATCAATGCTTTATCTCCTTGCCCCATAGTGC
CTAGAAAGTAACGGGGAAAGTAAATGCAAAGATAAAATTCTTGGCATTACAGCCTATGATACGACCCCTA
CCTTCTGCTAAGACATTGATGTGGATAAGGAAATTAGGTGCAACCAAGTGGATAAGGAAAATTATATCGATAGGG
GAAGAACCTTCTGGAAACAAGGCTCTGGTATGAATGTGGATATGCTCAGACCCATTGGGCAGAAAATTGCTAGT
GTGATGATGAAAATCAATGAGAAGCTAGGTGGCAAAGATTAG

(SEQ. ID. NO. 155)

MNLGEFWYNKINKRGRRLMKVRFIFLALLFLASPEGAMASDGTWQKQYLKEDGSQAANEWVFDTHYQSWFYIKADANYA
ENEWLKGDDYFYLKSGGYMAKSEWEDKGAFYLDQDGKMRNAWVGSYVGATGAKVIEDWVYDSOYDAWFYIKADGQHAE
KEWLQIKGKDYYFKSGGYLLTSQWINQAYVNAGAKVQQGWLFDKQYQSWFYIKENGNYADKEWI FENGHYYYLKSGGYMAAN
EWIWDEKESWFLKFDGKMAKEWVYDHSQAWYVFKSGGYMTANEWIWDKESWFLKSDGKIAEKEWVYDHSQAWYVFKSGG
YMTANEWIWDKESWFLKSDGKIAEKEWVYDHSQAWYVFKSGGYMAKNETVDGYQLGSDGKWLGGKTTNEAAYQVVPVTA
NVYDSDGKELSYISQGSVWLDKDRSKDDKLRAITISGLSGYMKTEDLQALDASKDFIPYYESDGHRFYHYVAQNASI PVASH
LSDMEVGKKYISADGLHFDFKLENPFLFKDLTEATNYSAEELDKVFSLLINNSLENKGATFKEAEHHYHINALYLLAHSA
LESNWGRSKIADKNNFFGITAYDTTPYLSAKTFDDVDKGILGATKWIKENYIDRGRTFLGNKASGMNEYASDPYWGEKIAS
VMMKINEKLGGKDZ

ID205 - 4118.1 (SEQ. ID. NO. 172)

ATGAAAAAAATTAGGTACATTACTCGTTCTCTTCTGCAATCATTCTGTAGCATGTGCTAGCGGAAAAAAAGATAACAC
TTCTGGTCAAAACTAAAAGTTGTTCTACAAACTCAATCATCGCTGATATTACTAAAATATTGCTGGTACAAAATTGACC
TTCATAGTAGTCTGTTCCGATTGGCAAGACCCACAGAATACGAACCACTTCTGAAAGACGTTAAGAAAATCTGAGGCTAAT
TTGATTCTTCTAAACGGTATTCAACCTTGAACAGGTGGCAATGCTGGTTTACAAAATTGTTGAGAAAATGCAAGAAAATCTGA
AAAACAAAGACTACTTCGAGTCAGCAGCGCTGATGTTATCTACCTTGAAAGGTCAAAATTGAAAGGAAAAGAACGACCCAC
ACGCTTGGCTAACCTGAAAACGGTATTATTGCTAAAATATGCCAACATTGAGGCCAAAGACCTAACAAATAAA
GAATTCTATGAAAAAAATCTCAAAGAATATACTGATAAGTTAGACAAACTGATAAAAGAAAGTAAGGATAAATTAAAGAT
CCCTGCTGAAAAGAAACTATTGTAACCAGCGAAGGAGCATTCAAATACTCTCTAAAGCCTATGGTGTCCAAGTGCCTTACA
TCTGGGAAATCAACTGAGAAGAAGAGAACCTCTGAAACAAATCAGGCTTGGTGAAGACACAAACATCCAACTACGCTCAAT
CTTACTGACTCTATCGAGAACAGGTAAGAAGGCGACAGCTACTACAGCATGAAATACAACCTTGACAAGATTGCTG
AAGGATTGGCAAATAA

(SEQ. ID. NO. 156)

MKKLGTLVLFLSAILVACASGKDDTTSQQLKVVATNSI IADITKNIAGDKIDLHSIVPIQDPHEYEPLPEDVKKTSEAN
LIFYNGINLETGGNAWFTKLVENAKTENKDYFAVSDGVDIYLEQNEKGKEDPHAWLNLENGIIIFAKNIAKQLSAKDPNNK
EFYEKNLKEYTDKLDKLDKESKDKFNKIPAEKKLIVTSEGAFKYFSKAYGVPSSAYIWEINTEEGTPEQIKTLVEKLROTKVP
SLFVESSVDRPMKTVSQDTNIPYAQIFTDSIAEQKGEQDSYSSMMKYNLDKIAEGLAKZ

ID206 - 4119.1 (SEQ. ID. NO. 173)

ATGGAATGGTATAAAAATCGGACTTCTGCAACTACAGGTTAGCTTGTGTTGGCTCGGGCTTGTCCAACATATGGTAA
ATCTCGGGATGGCACAGTGACCATCGAGTATTCAACCAGAAAAAGAAATGACCAAAACCTTGAAGAAATCACTCGTGATT
TTGAGAAGGAAAACCTAACAGATCAAGGTCAAAGTCGTCATGTACCAAAATGCTGGTGAAGTATTGAAGACACCGCGTTCTCGA
GGAGATGTCGCTGATGTGGTCAATATTACCCACAGTCCATCGAAGTGGCAAAAGCAGGTGTTGAAGATT
GAGCAACAAAGACTACTGAAACCGCTGAAAAATGGCTACCGTAAACGAAAATATGCTGTAACGAAAAGTTAACGTTCC
TTACAGCTAATGTTATGAAATTACTACAACAAAGATAAAATTGCAAGAACTGGCTTGAAGGTTCTGAAACCTGGATGAA
TTTGAACAGTTAGTCAAAGATATCGTGTCAAAGGACAAACACCATTGGAATTGCAAGGTGAGATGCTGGACACTCAATGG
TTAACATCAATTAGCCTTGCACAGCAACAGGGAGGAAAAGAAGAACATCAACCTTCGATCAATGATCTAACCAAATGCCA
TTAAATTGTCGATCGGATTGAAAGATGATCAAGGTGATGGACATCGGACATCCTCGCATGACATGATCTAACGAAAAGACTGG
GAAGGGTGTGGCTACCGATGTTATGGAGCCTTCGACAGTGGGATGCTCTGATCACACCAAATGGGCTTGGGGATCAC
AGCGATTAATGAAACAAAACGAACTTAAAGATTGGACCTCATGATTCCAGGAAAAGAAAAGGACAAAGCTTAACCGTTG
GTGCGGGAGACTTGGCATGGTCTATCTCAGCCACCACAAACATCCAAAAGAAGCCAATGCCCTTGTGAATATGACCGT
CCAGAAGTCATGCAAAACTACGATGTGGACGGATCTCCACAGCGATCGAAGGGTCAAAACAAAGCAGGAGAAGATTCA
GCTGCTGGTATGACCGAATATGCCCTTACGGATCGTACTGGTCTGGTCAACAAACTGGACAGTGAAGCAGACTTCC
ATACCTTGACCATGAACTATGTTGACCGGTGATAAAACAGGACATGGTCAATGATTGAATGCCCTTAAACCGATGAAA
GCGGATGTGGATTAG

(SEQ. ID. NO. 157)

MEWYKKIGLATTGLALFGLGACSНЫГKSADGTVTIEYFNQKKEMLTLEEITRDFEKENPKIKVVKVNPNAMEVLKTRVLA
GDVDPVVNIYPOSIELQEWAKAGVFDLSNKDYLKRVKNGYAEKYAVNEKVNPFNTANAYGIYNNKDKFEELGLKVPETWDE
FEQLVKDIVAKGQTPFGIAGADAWTLNGYNQLAFAATATGGGKEANQYLYRSQPNAIKLSDPIMKDDIKVMDILRINGSQKNW
EGAGYTDVIGAFARGDVLMTPNGSWAITAINEQKPNFKIGTFMIPGKEKGQSLTVGAGDLAWSISATTKHPKEANAFVEYMTR
PEVMQKYYDVGSPATAIEGVKQAGEDPLAGMTEYAFDTRHLVWLQQYWTSEADFHTLTMVNLQGMVNDLNAFFNPMK
ADVDZ

ID207 - 4123.1 (SEQ. ID. NO. 174)

ATGAAGAAAATCAAACCGCATGGACCGTTACCAAGTCAGACTCAGCTAGCTTATCTGGAGATGAACCTAGCAGCTTTATCCA
CTTCGGTCTCAAATACCTTTATGACCAAGAATGGGGACTGGACAGGAGGATCTGAGCGCTTAAACCGAGTCAGTTGATG
CGCGTGAAGTGGGTTCTGTCAGCAAGGAAAACGGGCTTAAAGGTTGATTGGTGTCAAGCACCACGATGGCTTGTCC
TATCCGACAGCTCACACAGATTATCGGTTAGGTCACTGCTTGGAGGAGAGGAAAAGGGCAGTGTCTCCTGAAAGTACCCCA

AGCTGCCACAGAGTTGATATGGATATGGGGTCTACCTGTACCGTGGGATGCCCATAGTCCCTCTATCATGTGGACCGAG
 AAGCGGACTACAATGCCATTATCTGGCTCAAGTGAAGGAAATCTTATCAAATCTAACTATGGGAATGCTGTAAGTCGCT
 GAGGTTGGATGGATGGTGCAGAGGAGAGGGCGCGAAAAGGTTAATTATGAATTGAAAAATGGTTGAAACCATTCTGTA
 CCTGCAGGGCGATTGCTGATTTCACAGAAGGACCCAGTATCCGCTGGATTGGCAATGAACGAGGGTATGCAAGTGATC
 CACTGTGGCAAAGGTTAATTCTGATAAAACTAGGAACAGAAGCAGAGCTGAACATCTCAGCACGGGATCCCTCGGGCAG
 ATTGTTCACTGGAGAGGAGCAGATGTTCCATCCGTCAGGCTGGTTCACTCATGAGGATCAGGATCTAAGTCTCGAGGA
 GTTGGTCGAAATCTACTTCACTAGAGGAGGAACTCCACTCTGCTTAATATTCCGCCAATCAAGCTGGCTTTG
 ATGCAAAGGATATTGAACGACTTTATGAATTGCGACCTATCGCAATGAGCTCTATAAAGAAGATTGGCTCTGGGAGCTGAG
 GTATCTGGTCCAGCTTCCGCAGACTTGTGCTGCATTGACAGACGGCCTTGAGACAGCTCTGGGCAAGCGATGC
 AGACTTGCCTATCCAGTAGAACTCGACTTAGGTTCTCTAAAACCTTGTGATGTAATTGAGTTAAGGAAAGATTGAGCTAG
 GGCACAGAACATGCTGCTTTCATGTGCAAGTAGAGGTTGGATCTGGCAGGAGTTGGTTGGGTCATACTGTTGGTTAC
 AACAGTCTCTACGAGGAGCAGTTGAGGACAGAAGATACGTGATGTCATTACAGAATCACAGGCTTGCCTTGTGAC
 CAAGATTCCCTTATAAAACTCCTGGATTATCAAAAAAAGAAGTTGTTAGGAACATGCAATTGAGGAAAGCCTAGCTG
 TGGCAAAGGAGAAAATGCCTATTTCAGTTAAGCGCAGAGAATGTAAGTGTGCTCTTAAAGCTAAGATTGAGCTAACC
 GGGACAGGTGTCCTAGGTGCGCTATCAGGATGAGATTCAAGTCCTTGCGTTCAACTGGTGAAGACTGAAAAAGTCTGAC
 GCTACCAACCTGTATTTCGAGGAGATAAAACCTGGATTATCTGAACCTACGGTGGATGGTCAGCTGTGGATCAAC
 TTCAGTCCAAGTTCAAA

(SEQ. ID. NO. 158)

MKKIKPHGPLPSQTLAYLGDELAIFIHFGPNTFYDQEWTGQEDPERFNPSQLDAREWVRVLKETGFKKLILVVKHHDGFV
 YPTAHTDYSVKSPWRRKGDLLEVSQAATEFDMDMGVYLPWDHSPLYHVDREADYNAYYLAQLKEILSNPNYGNAGKFA
 EVWMGDARGEAGOKVNVEFEKWFETIRDLOQDCLIFSTEGTSIRWIGNERGYAGDPLWQKVNPDKLGTEAELNYLQHGDPSGT
 IFSIGEADVSIRPGWFYHEDQDPKSLLELVEIYFHSVGRGTPLLNIPPNQAGLFDAKDIERLYEFATYRNELYKEDLALGAE
 VSGPALSADFACRHLTDGETSSWASDADLPQLELDLGSPKTFDVIELREDLKLQRRIAFHVQVEDGVWQEFSGHTVGY
 KRLLRGAVVEAQKIRVVITESQALPLTKISLYKTPGLSKKEVVQELAFAEKS LAVAKGENAYFTVKRRECSCGLEAKISIOP
 GTGVHVAYQDEIQVLAFTQTGETEKSLTLPFLYFAGDKTLDFYLNLTVDGQLQVQVSZ

ID208 - 4125.12 (SEQ. ID. NO. 175)

ATGCTTGAAAGACTGAAAAGAATACATTATGTTTGGATCAGTTAATTGATTTCCCCTCCTGCTGTAGTGAC
 TGGTGGCTTCTGCCTGGCATTTATGGTGTGATTCTATGTTAGGGTGTGTTAACAAACTAACAGGCCAGC
 GCCTATCTGGCTATATTGGGCTCATGCTGACTATGTTAGGGAAACTTGCCCTTGTGCTGTTAACAAACTATCTGGTT
 TTCTTTCTCATCCAATCTCTTAAAGTATCATTCAGCGTACGTAGTTAAAGTCTTACATGCTGACTTCTCTTC
 TCAAGTCTTGTGTTGGCAACTGTTGATTTTCAGAGAATCGAAGTTGAGTTCTATTCTATCTACTTGTAACTTCTACTT
 TTGTCGATTAACTGACTTTGGATTGGTTCGGATTGCTATTGCGAGGATTGAAAGAAGCTCAGGTCAGGATTTGCAAG
 ATAAATCTATGCTGAAATGACAGTGTGCTATCGTCAAGGATTTGCAAGTGTGACTCTGGACATACCTTGCTATGCT
 GAGGTGCAAGACAGATTAGCCTGCAAGTTCAAGATGGAGGTTATCCACAGGTTGGAAAAGGAATTAAAGAAATTCA
 AGATCAGCAAGGATCATGA

(SEQ. ID. NO. 159)

MLERLKRIHYMFWISLIFMIFPILSVTGWLSAWLLIDILFVVAYLGVLTTKSQRLSWLYWGLMLTYVVGNTAFVAVNYIW
 FFFLSNLLSYHFSVRSLSKSLHVWTFLLAQVLVVGQLLIFQRIEVEFLFYLVLTVFVLMTFGLVIRIRIVEDLKEAQVKQNAQ
 INLLAENERSRIGQDLHDSLGHFTAMLSVKTDLALQLFQMEAPQVEKELKEIHQISKDPZ

ID209 - 4126.3 (SEQ. ID. NO. 176)

ATGAATGATAAGTTAAAATCTTCTGCTAGGAGTATTGTTCTAGGCCATAACCGGTTCTATGTTCTATTGATACGAAA
 TGCAAGGGCAGACAGATGCCCTGCAAATGAAAGGCGGAGTTAGCCAAGGAGGAAAGCAGTGAACAAACAGGTCATGCAGAAC
 CGCCATGTCACTATGATGCAAGAAATTCTCTCAAGGAAATCTGCTTCACTATTCCGATTATCAATCCAACCA
 AAAATAAACAGGAAATGATGTTGGGACCCAGTACCGTACTGGTTATTACACAGATGACAAGGATTGGAAGTGAAC
 CAAGTCTTGTGAGGTTGCTAAAGAAATACGATCAACCTCTAGCAGTTGAAAGAAGGAAACTTGAAGAATTGTTGGTGGCTGA
 GGATTACCATCAAGACTATCTCAAGAAAATCCAATGGCTACTGCCATATCAATGTTAATCAGGCGCTATCTGTCATTG
 ATGCCAGCAAATATCCAAAAGCAGATTGAGGAATTGAAAAAGACCTGTCACCTGAGGAGTATGCAAGTTACCCAGGAAAT
 CAAACAGAACAGCTTCTCAAACCGTTACTGGGATAATTGAAATCCGTTATCTATGTTGAGATAGCAACTGGGAACCTCT
 CTTTCTATCAAAGACAAATTGAGTGTGGTTGGCTGGCTAGTTTACCCAAACCCATCAGGCCAGATGTTGTCACCTACA
 AGGAAGATAAGTCTACAAATGACCGTGTGGAGCTGGCAGAGTAGGAGATCTCACCTGGGATGTTGACGGAT
 GGTCCACAGGACAAGGGCGCTTACGTTACTGTATCAATAGCCTCTATCCGCTTATTCCAAAGACCAAATGGAAGAAA
 AggctACGCTTATTACTAGATTATGTTGATTAA

(SEQ. ID. NO. 160)

MNDKLKIFLLGVFFLAITGFYVLLIRNAGQTDASQIEKAASQGGKAVKTEISKDADLHEIYLAGGCFWGVEEYFSRVPG
 TDAVSGYANGRGETTKYELINQTHAETVHVTYDAQKISLKEILLHYFRIINPTSKNKQGNDVGTQYRTGVYYTDDKDLLEVIN
 QVFDEVAKKYDQPLAVEKENLKNFVVAEDYHQDYLKKNPNGYCHINVNQAAVPVIDASKYPKPSDEELKKTLSPEEYAVTQEN
 QTERAFSNRYWDKFESGIYVDIATGEPLFSSKDKFESGCGWPSFTQPISPDVVTYKEDKSYNMTRMEVSRVGDSHLGHVFTD
 GPQDKGGLRYCINSLSIRFIPKDQMEEKGAYLLDYVDZ

ID210 - 4127.1 (SEQ. ID. NO. 177)

ATGAAAAGAAAATGGATGATTATGCTGCTTCTATGAATCTGCCATGACAGTTCTGATAAAGGAGACGGCG
 TCGCTAGTCAGCTTATTCAACAAACTCAGAGGGCTTAATTGGAGCAACTATTCTGCCCTTGAGAAAATATGGTATCAAAG
 TAGAATGATCAAGCTGGTACTGGAGAACTTTCAAAAACACTAGAGTCAGAAAAGAAGTCTGCTAGCTGATGTTATCTTT
 GGTGGTTCTTATACAAATATACTACCCACGGAGAACTCTTGAAAAACACTATCTAAAAGAAAATGATAATGTTATCAAAGA
 ATATCAAACACAACGGCTACTCTACTCCTTATACACTAGATGGTAGTTAACCTGTAACCCCTGATTAACTAAAGGCA
 TGAACATCGAAGGATAAACGATCTTCAACACTGAACAAAAGGAAAATCGCAACTGCTGACCCAGCAAACCTCTAGC
 GCCCTTGCTCAATTAACAAATATGCTACAAGCTCAAGGTGGTACAAAGATGATAAGGCTTGGCTTATGTTAAAGATCTTT
 CACACTTATTGATGGTAAATCGGTTCAAGTTCATCTAGTGTCTATAAAGTAGTGTCTGATGGAGAAATGGCTTGGCTCT
 CTTATGAAGATCAGCAGTAAACTCTTAAATGACGGAGCTAACATTAAGGTTAGTGTCTATCCAAAAGAAGGAAACCGTCTC
 CCTGCTAGTGTCTATCGTTAAAATCTAAAGGAAATTGCAAGAAATTATGCAAGTATTATCTCTCAAGAAGT

ACAAGATACACTTGGTACAACCACTACTAACCGCCTGTCGAAAAATGCTAAAACAAGCAGAAACATGAAACCAATTGACA
AAATCAAACACTCACTGAAGATTATGATTATGTCATCAAGAATAAAATCAGATATCGTTAGAAATACAACGAAGTCTTACA
GATATCCAATCTAAACAGTAA

(SEQ. ID. NO. 161)

MKKKWMYYAACSSNESADDSSDKDGGSLLVVYSPNSEGGLIGATIPAFEEKYGIKVELIQAGTGELFKKLESEKEVPVADVIF
GGSYTQYTTHGELFENYTSKENDNVKEYQNTTGYSTPYTLGDSLIVNPDLTKGMNIEGYNDLFKPELKGIATADPANSS
AFAQLTNMLQAQGGYKDKAWSYVKDLFTLIDKGKIGSSSSVYKVVADGEMAVGLSYEDPAVKLNDGANIKVVYPKEGTVFL
PASAAIVVKSKNMENAKKFIDFIISQEVDLGTNTNRPVRAKNTSENMKPIDKIKTLTEDYDYVIKNKSDIVKKYNEVFT
DIQSKQZ

ID211 - 4127.2 (SEQ. ID. NO. 178)

ATGAGTGAGATCAAATTATTAACGCCAAAAAAATCTACCACGATGTCCTGTTATTGAGAATTGAAACATTACAATTCCAAA
AGGAAGTCTCTTACCCCTCTGGAGCTTCAGGATGTTGGAAAACGACCCCTCTTCGATGATTGCAGGTTCAACAGTATCG
AAGGAGGAGAATTACTTCGATGATAACAAATATGGAACACGAGCAGGAAACGCAATATCGGGATGTTTCCAAAAC
TACGCTATTTCCTCACATTGACTGTGAGGACACGTTGCTTTGGCTTATGCAAAAGGTTCCAAAAGAAGAATTGAT
TCAACAGACCAACAAGTATCTGAACATGCAAATTGCTAATATGGGATCGAAAGGCCATAACTCAGTGGTGGACAC
AACAACTGTGACCTTGGCATGCGCCTAGCGGTTAATCCAAGTGTCTCTCATGGACGCCACTTAGTAATCTGGAGGCC
AAACTTCGCTGGATATGCGTCAAGGCCATCCGAGAAATCCAAACACGAAGTGGGATACAAACTGTTTATGTAACCCACGACCA
AGAAGAAGGCCATGGCTATTTCGACACAAATTGCTGTTATGAGAAGTGGGATGCAACAAATATCGGCCGACCAAAAGAAGTCT
ATCATAAAACAGCTATGAGTTGCGCAACCTTATCGGAGCAGCACAATTATTCCTGCAATCTGAAAAACGGAGGAC
GGCGCTTATATCGTCTTTCAGATGGCTATGCCCTCGAATGCCAGCTTGATCAGGTTGAGGAGCAAGCTATTGATGAA
CATTGTCAGAGAGTTTATCAAAGATGAATCTGGAGATATTGAAGGAACATTAGAGATAGCTTATCTGGACTAAATA
CGGATTATTCTCATGAGACAGGTTTGCCTCAAAAATTCAAGTTAGTGAAGAATCAACTTTGAAGAAGATCTACAAAAGGC
AATCGTATTGCTACGAATCAATACGCAAAATTAAACATCTTCTCAGATGGTCCAAAACCTGATAAAAGGAGTCAA
CCATGGAACGTA

(SEQ. ID. NO. 162)

MSEIKIINAKIYHDVPIEVLNITPKGSLFTLLGASCGKTLRLMIAGFNSIEGGEFYFDTKINNMEP SKRNIGMFQN
YAIFPHLTVRDNVAFGLMQKKVPKEELIQQTNKYLELMQIAQYADRKPDKLSSGQQQRVTLACALAVNPSVLLMDEPLSNLEA
KLRDLMRQAIREI QHEVGITTVYVTHDQEEAMAISDQIAVMKDGVIQQIGRPKELYHKPANEFVATFIGRTNIIPANLEKRSD
GAYIVFSDGYALRMPALDOVEEQAIHVSIRPEEFIKDESGDIEGTIRD SVY LGLNTDYFIETGFASKIQVSEESTFEEDLQKG
NRIRLRINTQKLNIFSADGSQNLIGVNHGTZ

ID212 - 4136.1 (SEQ. ID. NO. 179)

ATGAAGAAAAATTATTGGCAGGTGCCATCACACTATTATCAGTAGCAACTTTAGCAGCTTGTGAAAGGGTCAGAAGGTGC
AGACCTTATCAGCATGAAAGGGATGTCATTACAGAACATCAATTATGAGCAAGTGAACAGCAACCCCTCAGCCCACAAAG
TCTTGTAAATATGACCATCCAAAAGTTTGGAAAACATGGCTCAGAGCTTGTGATAAAAGAGGTTGATGATACTATT
GCCGAAGAAAAACAAATATGGCAGAAACTACCAACGTCTTGTCACAAGCAGGATGACTCTGAAACACGTAAGCTCA
AATTCTGCTACAAGTAAATTAGTTGAGTTGGCAGTTAACATCGCTTAATATGAGATAAGGCAAAAGAAGTCTCGAAAAGCCT
TTGATGAGTACACTCCAGATGTAACGGCTCAATCATCGCTTAATATGAGATAAGGCAAAAGAAGTCTCGAAAAGCCT
AAGGCAGAAGGTGCTGATTTGCTCAATTAGCCAAAGATAATTCAACTGATGAAAAACAAAAAGAAAATGGGAGAAAATTAC
CTTGATTCTGCTTCAACAGAAGTACCTGAGCAAGTCAAAAAGCCCTTCGCTTATGAGTGGATGGTGTGATGTGA
TTACAGCAACTGGCACACAGCCTAACAGTAGCCAAATTACATTGTAACACTCACTAAGAAAACAGAAAATCATCTAATATT
GATGACTACAAAGAAAATTAAACTGTTATCTGACTCAAAAACAAAATGATTCAACATTGTTCAAAAGCATTATCGGAA
AGAATTGCAAGCAGCAATCAAGGTTAGGACCAAGCCTCCAAAATATCTTACCCAAATATCGGTGGAGATTCAA
GCTCAAGCAGTAGTACATCAAACGAATAG

(SEQ. ID. NO. 163)

MKKKLLAGAITLELSVATLAACSKSEGADLISMKGDVITEHQFYEQVKSNSPAQQVLLNMTI QVFEKQYGS ELDKEVDDTI
AEEKKQYGENYQRVLSQAGMTLETRKAQIRTSKVELAVKVAEAELTDEAYKKAFDEYTPDVTAQIIRLN NedAKEVLEKA
KAEGADFAQLAKDNSTDEKTENGGEITFDSA STEVPEQVKKAFA L DVGSDVITATGTOQYSSQYYIVKLTKKTEKSSNI
DDYKEKLKTVILTQKQNDSTFVQSII GKE LQAA NI KVKDQAFQNIFTQYIGGDSSSSSTSNEZ

ID213 - 4137.3 (SEQ. ID. NO. 180)

ATGAAAAAAATTAACAAATATGTAACCTTAGGTACTGTAGTGGTATTATCAGCATTGTCGTAACTCAGTGCAGCTCA
GGAGACTGAAACTTCTGAGTCAACACCAAAGTGGTCAACCTGTCACCAAGCAGCTCCGATTCGGAAAGTACAACCTA
CATCGGATAACTCTCGGAAGTTACTGTCACACCTCGAACAGTGAACACTACTGTTAAGGATCCATCTCTACAGCGGAAGAA
ACTCTGTCTAGAAAAAAATATGTTACTTAAACAGGGGGCGAGAAAATGTTACTAAGAGTTAAAGGATAAAATTACTAG
CGGTGACTTTACTGTAGTATTAGTACAATCAGTCAGTGAGAAAGGTTACAAGCTCTGTTGGATATCTAATTCCAAAC
CCGGTCAACAAATAGTTAGTGTAGTTGGCTTAGAGACATGGTAGTTGGGATGGAAGCGCGTGATACTCTTCCAA
AAAAAAATACCTAGTATCCAGACCTGCTTCAGTTGGGTAAGTACAAACAGAGGCTGTGACTAACACTGTTGAGTAG
AGATTCACTGCAAAAAACATATTCTTACGCAAAATGGTACAAAAGTAGTAGAAAAGAAAAGTGGATAATTCTAAACATCA
AGGATATTAAAGGTTAGTACTATATGCTTGGGGAGTGAAACGTCAGGAAAACGGCTTGGTTAACGGAAACACTA
GAAAATATCAAATTCTTAATAGTGATTGGATGAAAGAAACTGTTAAAAGATGACAACAAACGCTGTTACTGGACATTAAAT
TTATACGGCTAATGATAACACAGGTTCTAATTTCCGTATTCCAGTTCTGTATACTTTAGCAATGGTCGGGTATTCTCAA
GCATTGACGCTCGTTACGGTGGAAACTCATGATTCTGAAATAAAATTAAATGCTACAAGTTAGTGTGATAATGGTAAG
ACATGGACTAAACCAAAAATTAAACATTGGCATTGATGATTTCGCGCAGTACCAATTAGAATGGCTCGTGAAGTTGGGACG
TGACTTACAAATCAGCGTGGTGAACCTATATTGACTCTGTTATTGTTAAAACAAATTGATGGTAAATTACCTT
AAATTAAAGGAAACAAGGTGATACTGATTACAATTACTATTCGTGAGAATGGTACTGTATACGACGATCGTACCAACAGACC
AACTGAATTTCAGTAGATAAAAATTGGTATTAAACAAATGGTAAATTGGTACGGTAGAGCGG

(SEQ. ID. NO. 164)

MKKNIKOYVTLGTVVVLSAFVANSVAQETETSEVSTPKLVQPVAPTPISEVQPTSDNSSEVTVQPRTVETTVKDPSSSTAEE
TPVLEKNNVLTGGGENVTKELDKFTSGDFTVVIKYNQSSEKGLQALFGISNSKPGQQNSYVDVFLRDNGELGMEARDTSSN
KNNLVSRPASVGKYKQEAVTNTVAVVADSVKKTYSLYANGTKVVEKVDNFNIKDIKGIDYYMLGGVKRAGKTAFGFNGTL
ENIKFFNSALDEETVKMNTNAVTGHLIYTANDTTGSNYFRIPVLYTFNSGRVFSSIDARYGGHTDFLNKINIATSYDDNGK
TWTKPKLTLAFDDFAPVPLEPREVGRDLQISGGATYIDSVIVEKKNKQVLMFADVMAGVSREATRKDSGYKQIDGNYYL
KLRKGQDTDYNYTIRENGTYDDRTNRPTEFSVDKNGFIKQNGNYLTVER

ID214 - 4185 (SEQ. ID. NO. 181)

ATGAAAAAAATTAGCCTATTACTAGCTATCCTACCATTTTGTTGCCGTGAGAATCAAGCTACACCCAAAGAGACTAGCGC
TCAGAAAGACAATCGCCTTGCTACAGCTGGCGACGTGCCACCATTGACTACGAAGACAAGGGCAATCTGACAGGCTTGATA
TCGAAGTTAAAGGCACTAGATGAAAACACTAGCGACTACGAGATTCAATTCAAAGAACCGCCTGGGAGAGCATCTTCCA
GGACTTGATTCTGGTCACTATCAGGCTCGGGCCAATAACTTGAGTTACACAAAAGAGCGTGTGAAAAATACCTTACTCGCT
TCCAATTCCAACAATCCCCTGCTTGTCAAGCAACAAGAAAATCCTTGACTTCTCTTGACCAGATCGTGTGAAACAA
CACAAAGAGGATACCGGAACTCTAAGCTCAATTCATCAATACTGGAACTCAGAAACACACTGATAATCCGCTACAATTAAAT
TTTCTGGTGGAGGATATTGGTAAAGAATCTAGACCTTGCTAACGGAGAGTTGATTCTCTAGTTTGACAAGGTATCCGT
TCAAAGATTATCAAGGACCGTGGTTAGACCTCTCAGTCGTTGATTACCTTCTCAGATAGCCCCAGCAATTATATCATT
TCTCAAGCGACCAAAAGAGTTAAAGAGCAATTGATAAAGCGCTCAAAGAACTCTATCAAGACGGAACCTTGAAAACCT
AGCAATACCTATCTAGTGTTCTACCTCCAGATCAATCTCAGTTACAATAA

(SEQ. ID. NO. 165)

MKFKSLLAILPFLVACENQATPKETSQAQKTIVLATAGDVPPFDYEDKGNLTGFDIEVLKAVDEKLDYIEQFORTAWESIFP
GLDSGHYQAAANNLNSYTKERAEKYLSPNPLVLSNNPLTSNPLDQIAKGTTTOEDTGTNSAOFINNWOKHTDNPATIN
FSGEDIGKRILDLANGEFDLVDKVSQKIIKDRGLDLSVVDLPSADSPSNYIIFSSDQKEFKEQFDKALKELYQDGTLKEL
SNTYLGGSYLPDQSQLQZ

ID215 - 4211.1 (SEQ. ID. NO. 182)

ATGAAAAAAATAGTTATATCATCCTCACTCTTTTGCTTGTTCTTATTGTCTATGCTACGGCGACGAATTTC
AAACAGTACCAAGTGCTAGGCAGGTAACCGAAACCTATACTAATACAGTAACAAATGTCCTATTGACATACGCTATAATA
GTGATAAGTATTTATTAGCGGTTTGCTTCAGAAGTATCAGTGGCTTGACTGGTCAAATCGCCTATCGTAGCTAGTGA
ATGCAAGAAAGTACACGTAAGGTTACTGCTGACCTAACAGATGCCGTGTTGAAAGTTGAGTTCCCTTGAGCAT
TGAAGATTACCCAATGGGCTGACCGCTGTGGCAGCTCCGCAAAACATTACAGTCAAGATTGGTAAGAAGGCTCAGAAGGATA
AGTAAAGATGTACCAAGAGATTGACCTAGTCACATTGATAGTCGGGTACAATTGAAAATGTCATGGTGTGAGATAAAAGAA
GTGTCTATTACGAGTGACCAAGAGACATTGGATAGAATTGATAAGATTACGCTGTTTGCCAAGTAGCGAACGTATAACAGG
TAATTACAGTGGTTAGTACCTTGCAAGGCAATCGACCGAATGGTGTGTTACCGCAGTTACACTCGTTGATACAA
TAATGAAGGTGACTACAAAACCAGTAGCACCAAGTTCAAGCACATCAAATTCAAGTACAAGCAGTTCATCGGAGACATCTCG
TCAACGAAAGCAACTAGTTCAAAAACGAATTAA

(SEQ. ID. NO. 166)

MKNSLYIISLFFACVLFVYATATNFQNSTSARQVKTETYNTVTNPIDIRYNSDKYFISGFASEVSVVLGANRLSLASE
MQESTRKFKVADLTDAGVGTIEVPLSIEDLPNGLTAVALPQKITVKIGKKAQKDKVKIVPEIDPSQIDSRVQIENVMVSDKE
VSITSDQETLDRIDKIIAVLPTSERITGNYSGSVPLQAIIDRNGVVLPAVITPDFTIMKVTTKPVAPSSSTSNSSETSS
STKATSSKTNZ

ID216 - 4127.3 (SEQ. ID. NO. 183)

ATGTTGATTGGCGAAGGGTATCGGACTTCCCTGCTGATTTATACCCAATTAGCGAGGTTGGAGGAAATTCTGCTTT
TGCATTATGGCATTATCATTGCTTGGCAATTTCCTTATCCAAAACACATTGCAAACCGCTACAGTTCAGCATGAATC
TGCTCCATCCAATTGAGCCTAAAAAAACTACAAAGGAAAATGGCTGCCATTATGCAACAGTCTACGGAATTATCTTATC
TCTGTTTACCTCAAATCTACTAATTATACCTCTTCCCTAAAACATCAGGTATGGTATCTGTTAAAGGTATTCTCAA
CAGTTACAAGGTAGCTTCCATCGTAGGGATCTGCTATTTCATACCCATTGCTATCCCTTGATTGCTTAGTTCTAGTTG
TTCTATTGCGACATTATCTCTACCTAGGGCTTAGGGATTGCGCTTATTCCTTCAACTACTGGTCTATTGCAAGCCTCAGTATGGTA
CCTTATATTGCGACATTGCGCTTAGGGATTGCGCTTATTCCTTCAACTACTGGTCTATTGCAAGCCTCAGTATGGTA
GATTACAGGACTGCTTCTGATTATGTCCTATCTGCCAGAAGATTACCTTACTATTGCTCATCTGCTTAAGATTACA
TACAACAAATAGCACCAAGTATTGAAGAAGCTGCTGAAAGCTTAGGAAGTAGTCGCTCAATACCTTGCTAAGATTACA
CCAATGATGCTATCTGGTATCATTCTGGAGCCATCTTGA

(SEQ. ID. NO. 167)

MLIGEGYRTFPVLIYTOFISEVGGNSAFAIMAIIALIIFIQKHIANRYSFSMNLLHPIEPKTTKGKMAIYATVYGIIFI
SVPQIYLIYTSFLKTSGMSVKGYSPNSYKVAFHRMGSIAFNTRIRIPLIALVVLFATFISYLAVRKRNLFNLIDSLSMV
PYIVPGTVLGIAFISSFNTGLFGSGFLMITGTAFILEIMSLSARRLPYTIRSSVASLQIAPSIEEAAESLGSRLNTFAKITT
PMMLSGIISGAILSZ

Table 4

ID301 (SEQ. ID. NO. 196)

ATGAATAAGAAAAAATGATTTAACAGTCTAGCCAGCGTCGTATCTTAGGGCTGGTTGTTACGTCAGCCTACTTT
 TGTAAGAGCAGAAGAATCTCCAAGTGTGCAAAATCTCATTAGAGAAGAAATATGAGGAAGCAGAAAAGCTGATA
 CTGCCAAGAAAGATTACGAAACGGCTAAAAGAAAGCAGAAGACGCTCAGAAAAGATGAAGATGATCAGAAGAGAAACTGAG
 GAGAAAGCTCGAAAAGAACAGAAGCATCTCAAAAATTGAATGATGTGGCCTGTTGTTCAAAATGCATAAAAGAGTACCG
 AGAAGTTCAAAATCAACGTAGTAAATATAATCTGACGCTGAATATCAGAAAAAATTAACAGAGGTCGACTCTAAATAGAGA
 AGGCTAGGAAGAGCAACAGGACTTGCAAAATAAATTAAATGAAGTAAGAGCAGTTGAGTCTCTGAACCAAATGCGTGGCT
 GAGACTAAGAAAAAGCAGAAGAGCTAAAGCAGAAGAAAAAGTAGCTAAGGAGAAAATATGATTATGCAACTCTAAAGGAGTGC
 ACTAGCAGAAGAAAAGTAGAGGCTAAGGAACCTGAAATTGAAAAACTCTCAATATGAAATTCTACTTTGGAACAAGAAGTTG
 CTACTGCTCACATCAAGTAGATAATTGAAAAACTCTGCTGGTGCAGATCTGATGATGGCACAGAAGTTATAGAGCT
 AAATTAAAAAAAGGAGAAGCTGAGCTAAACGCTAAACAAGCTGAGTTAGCAAAAAAACAAACAGAACTTGAACCTCTTGA
 CAGCCTTGATCCTGAAGGTAAGACTCAGGATGAATTAGATAAAGAAGCAGAAGAAGCTGAGTTGAGTAAAGAGCTGATGAAC
 TTCAAAATAAAGTGTGATTTAGAAAAAGAAATTAGTAACTCTGAAATATTACTTGGAGGGCTGATCTGAGATGATACT
 GCTGCTCTCAAAATAAATTAGTGTGCAAAAGCTGAGTTAGCAAAAAAACAAACAGAACTTGAACAAACTCTTGACAGCCT
 TGATCTGAGGTAAGACTCAGGATGAATTAGATAAAGAAGCAGAAGAAGCTGAGTTGAGTAAAGAGCTGATGAACCTCAA
 ATAAAGTTGTGATTTAGAAAAAGAAATTAGTAACCTGAAATATTACTTGGAGGGCTGATTGAGTAAAGATGATACTGCTGCT
 CTTCAAAATAAATTAGTACTAAAAAGCTGAATTGAAAAAAACTCAGAAAGAATTAGATGAGCTCTTAATGAGTTAGGCC
 TGATGGAGATGAGAAGAAGAACTCCAGCGCCGCTCCTCAACAGAGCAGACACCAGCTCTGCACAAAACCAGAGCAACCAGCTC
 CAGCTCCAAAACCAGAGCAACCAGCTCTGCACAAAACCAGAGCAGACACCAGAGCAACCAGCTCCAAAACCAGAGCAACCAGCTC
 GCTCCAAAACCAGAGCAACCAGCTAGCCGGAGAAACCAGCTGAAGAGCTACTCAACAGAGCAGCTCCAAAACCAGAGCAGC
 AGCTGGAAACAAGAAAACGGTATGTTGTTACCTACAATACTGATGGTCAATGGCAATAGGTTGGCTCAAACACGGTT
 CATGGTACTACCTAAACGCTAACGGCGTATGGCAACAGGTTGGTGAAGAGATGGAGATACCTGTTACTATCTGAGCAGATCA
 GGTGCTATGAGAAGCAGCAACGGTCAAAAGTATCAGATAAATGGTACTATGTCACAGCAATGGCGTATGGCGACAGGGCTG
 GCTCCAATACAATGGTCTATGGTACTACCTCAACGCTAAGTGTGATATGGCGACAGGATGGCTCAAATACAACGGTTCTGTT
 ATTACCTCAACGCTAATGGTGTATGGCGACAGGATGGCTCAAAGTCAACGGTTCTGTTACTACCTAAACGCTAACGGTCTGTT
 ATGGCTACAGGTTGGCTAAAGTCAACGGTTCTGTTACTACCTAAACGCTAACGGTTCAATGGCAACAGGTTGGTGAAGA
 TGAGAGATACCTGGTACTATCTTGAAGCATCAGGTGCTATGAAAGCAAGCAATGGTCAAAGTATCAGATAATGGTACTATG
 TCAATGGCTTAGGTGCCCTGAGTCAACACAACGTAGATGGCTATAAGTCAATGGCAATGGTGAATGGTTAA

(SEQ. ID. NO. 184)

MNKKKMLTSLASVAILGAGFVTSQPTFVRAEESPQVVEKSSLEKKYEAKAKADTAKKDYETAKKKAEDAQKKYEDDOKRTE
 EKARKEAEASQKLNDVALVVQNAYKEYREVQNQRSKYKSDEYQKKTTEVDISKIEKARKEQQDLQNKFNEVRVVPEPNALA
 ETKKKAEEAKAEVKRKYDYLTKVALAKKEVEAKELEIEKLQYEISTLEQEVAQAHQVNDLKKLAGADPDDGTEVIEA
 KLKKGEAELNAQKQELAKKQTELEKLLDSDLPEGKTQDELDKEAEEAELDKKADELQNKVADLEKEISNLEILLGGADPEDDTAA
 AALQNLKLAAKKAEELAKKQTELEKLLDSDLPEGKTQDELDKEAEEAELDKKADELQNKVADLEKEISNLEILLGGADSEDDTAA
 LQNLATKKAELEKTQKELDAALNELGPDGDEEETPAPAPQPEQPAPAPKPEQPAQPKPEQPAQPKPEQPAQPKPEQPA
 APKPEQPAKPEKPAEERPTOPEKPATPKTGKQENGWMWFYNTDGSMAIGWLQNNGSWYLNANGAMATGWVKDGDWTYYLEAS
 GAMKASQWFKVSDKWYVNSNGAMATGWLQYNGSWYLNANGDMATGWLQYNGSWYLNANGDMATGWAKVNGSWYLNANGA
 MATGWAKVNGSWYLNANGSMATGWVKSDKWYVNGLALAVNTTDGYKVNANGEWZ

ID302 (SEQ. ID. NO. 197)

ATGTTTGCATCAAAAGCAGAAGAAAAGTACATTTCATTCAATTCTGAAATTAGTGTGGAGTAGCTAGTGTAGTTGCCAG
 TCTGGTATGGAGATGTGGTTCATGCCAGAGAAGCAGGGAGCTACCAAGTACCTCTCTCAATAGGGCAAATGAAA
 GTCAGGCAGAACAGGAGAACACCTAAAAACTCGATTCAAACGAGATAAGGCAAGGAAAGGGTCGAGGAATATGAAAA
 AAAATAGTGGGTGAGAGCTATGCAAAATCAACTAAAAGCAGACATAAATTACTGTTAGCTCTAGTTAACGAGTTGAACACAT
 TAAGAACGAGTATTGAATAAAATAGTTGAATCAACCTCAGAAAGCCAACAGATACTGATGATGGAGAGTCGATCAAAG
 TAGATGGAGCTGTGCTAAGTTGAAAAGGACTCATCTCTCGTCAGACTCTTCACTAAACCGGAAGCTCAGAT
 ACAGCGAACGCAACACGGCAGAGAACCCAGGAGAAAAGTAGCAGAGCTAAGAAGAAGGTTGAGAAGCTGAGAAAAAGC
 CAAGGATCAAAAGAAGAGATCGCTGTAACTACCCAACTTACATACAAACGCTGAGATTGCTGAGTCCGAT
 TGAAGTTAAAAGCGGAGCTGAACTAGTAAAGCTAACGAACCTCGAGACGAGCAGAAAAATTAAAGCAAGCAGAA
 CGCGAAGTTGAGAGTAAACAAGCTGAGGCTACAAGGTTAAAAAAATCAAGACAGATCGTAAGAAGCAGAAGAAGCTAA
 ACCAACAGCAGTCAAAGAGCAAGGTAACCAAGGGCGGCAAACAGAGGAGTTCTGGAGAGCTAGCAACACCTGATA
 AAAAGAAAATGATGCGAAGTCTTCAGATTCTAGCTAGGTGAAGAAACTCTCCAAGGCTCCATCTGAAACCCAGAAAAAG
 GTAGCAGAAGCTGAGAAGAGGTTGAGAAGCTAAGAAAAGCAGGAGATCAAAGAAGAGATCGCGTAACTACCCAA
 CAATACITACAAACGCTGAACTGAAATTGCTGAGTCCGATGTGGAAGTTAAAAGCGGAGCTGAACTAGTAAAAGAGG
 AAGCTAAGGAACCTCGAACAGGAAAAAGTTAAGCAAGCAAAGCGGAAGTTGAGAGTAAAAAGCTGAGGCTACAAGGTTA
 GAAAAAAATCAAGACAGATCGTAAAAAGCAGAAGAAGCTAAACGAAAAGCAGCAGAGAAGATAAAAGTTAAAGAAAAAC
 AGCTGAACACCACAAACCGCGCCGCTCCAAAAGCAGCTCCAGCTCCTCAAACCCAGAGAATTCAGCTGAACACAA
 CAAAAGCAGAAAACCGACTGATCAAAGCTGAGAAGACTATGCTGAGATCAGAAGAAGATAATACTCGCTTACTCAA
 CAGCAACCGCCTAAACGCAACCATCTACTCCAAAACAGGCTGAAACAAGAAAACGGTATGTGGTACTT
 CTACAATACTGATGGTTCAATGGCGACAGGATGGCTCAAACATGGCTCATGGTACTACCTCAACAGCAATGGCGTATGG
 CGACAGGATGGCTCCAAAACAATGGTCACTGGTACTATCTAAACGCTAATGGTCAATGGCAACAGGATGGCTCCAAAACAAT
 GGTCTATGGTACTACCTAAACGCTAATGGTCAATGGCGACAGGATGGCTCAAATCAATGGCTCATGGTACTACCTAAACGCT
 TAAATGGTCAATGGCGACAGGATGGCTCCAAATACAATGGCTCATGGTACTACCTAAACGCTAATGGTGTATGGCGACAGGTT
 GGGTGAAGAGTGGAGATACCTGGTACTATCTTGAAGCATCAGGTGCTATGAAAGCAAGCAATGGTCAAAGTATCAGATAAA
 TGGTACTATGTCATGGCTCAGGTGCCCTGAGTCAACACAACGTAGATGGCTATGGAGTCATGCAATGCAATGGTGAATGGGT
 AAACCTAA

(SEQ. ID. NO. 185)

MFASKSERKVHYSIRKFSGVVASVVASLVMGSVHATENEGATOVPSSRNANESOAEQGEOPKKLDSERDKARKEVEYVK
 KIVGESYAKSTKKRHTITVALVNELNINKEYLNKIVESTSESQQLILMMESRSKVDFAVSKFEKDSSSSSSDSSTKPEASD
 TAKPNKPTEPGEKVAEAKKVEEAEKKAQKDKQKEEDRRNYPITIYKTLLEIAESDVEVKKAELELVKVKANEPRDEQKIKQAE
 AEVESKQAEATRLKKIKTDRREEAEKKRADAKEQGKPKGRAKRGVPGELETDPDKENDAKSSDSVGEETLPSPSLKPEKK
 VAAEAKKVEAKKAEDQKEEDRRNYPITNTYKTLLELEIAESDVEVKKAELELVKEEAEPRNEEVKVKQAKAEVESKKAETRL
 EKIKTDRKKAAEAKRKAEEEDKVKEKPAEQPQPAKPAEKPAEPQPAEKPADQQAEEEDYARRSEEYNRLTQ

QOPPKTEKPAQPSTPKTGKQENGWYFYNTDGSMATGWLQNNGSWYLNNSNAMATGWLQNNGSWYLNANGSMATGWLQNN
GSWYLYNANGSMATGWLQYNGSWYLNANGSMATGWLQYNGSWYLNANGDMATGWVKDGTWYYLEASGAMKASQWFVSDK
WYVNGSGALAVNTTVGDGYGVNANGEVNZ

ID303 (SEQ. ID. NO. 198)

ATGGTAAAAAGACGTATAAGGAGAGGGACGAGAGAACCTGAAAAAGTTGTTCTGAGCAATCATCTATTCCCGTATCC
TGTATCTGTTACATCTAACCAAGGAACAGATGTAGCAGTAGAACAGCTAAAGCAGTTGCTCAAACAAACAGACTGGAAACAAG
AAAATGGTATGTTATTTTATAACTGATGGTCCATGGCAACAGGGTACAAGTTAATAGTCATGGTACTACCTC
AACAGCAACGGTCTATGAAAGTCATCAATGGTCCAAGTTGGTAAATGGTATTATGTAATACATGGGTGAGTTAGC
GGTCAATACAAGTATAGATGGTATAGAGTCATGATAATGGTGAATGGTGCCTAA

(SEQ. ID. NO. 186)

MVKRRIRRGTRPEKVVVPEQSSIPSYPVSVTSNQGTDVAVEPAKAVAPTTDWKQENGWYFYNTDGSMATGWWQVNSSWYLN
NSNGSMKVNWQFQVGGWYYVNTSGELAVNTSIDGYRVNDNGEWRZ

ID304 (SEQ. ID. NO. 199)

CTGAATACAAGTTGTCATGCTGCTGATGGGATTCAATATGTCAGAGATGATACTAGAGATAAAGAAGAGGGAAATAGAGTA
TGATGACGCTGACAATGGGATATTATTGAAAAGTAGCAGTAAACCTAAGGTAGTAACCAAGAAAATTCAAGTACCGC
TTCGTTATGAAAAGATGAAACAAAGACCGTAGTGGAAATCTGTTACAATTGATGGAGAGGATGGCTATGTAACACGACA
AGGACCTACGATGTTAACCGAGACTGGTACCGAACAGGGTACTGTTGATAGAAAAGAAGCCACGGATAACAGTTAT
CAAAGTTCCAGCTAAAGCAAGGGTGAAGAGTCTTGTGTTACTAATGTTGATAGAAAAGAAGCCACGGATAACAGTTAT
GACAGGAGCAAGAGATTACTCTAGGAAAAGATGGAAAACAGTTACACGATAACTTATAATGATGGAAAGAGTGGACAA
GTAACGAGAGTACTTTAAGTCAAAAAAAGACTCTAAACAGAGTTGTTAAAAAAAGAACCAAGCCCCAAGTTCTGCTCA
AGAAATTCCAATCGAAACAGAATATCTGATGCCCAACTCTTGATAAAAGTCAGAAGTAGAAGAGTAGGAGAAATTGGTA
AATTACTCTTACTACAATCTATACTGTAG

(SEQ. ID. NO. 187)

LNTSFVHAADGIQYVRDDTRDKEEGIEYDDADNGDIIVKVATKPKVVTKISSTRIRYEKDETDRSENPVTIDGEDGYVTTT
RTYDVNPETGYVTEQVTVDRKEATDTVIKVPAKSKEEVLPVFATKYEADNLDLSAGQEIEITLGKNGKTWTITYNVDGKSGQ
VTESTLSQKKSQTRVVKRTPQVLVQEIPETEYLDGPTLDSQEVVEVGEIGKLLLQSILZ

ID305 (SEQ. ID. NO. 200)

ATGAAGCTTTGAAAAAAATGATGCAAATCGCACTAGCCACATTTTCTCGTTTAGCGACAAATACAGTATTCAGA
TGATTCTGAAGGATGGCAGTTGTCAGAAGACCTACTACAAAAGGGGATCTAAAGAAAACCTACTGGAGAG
TGATAGATGGAAGTACTATTATTGATCCTTATCCGGAGAGATGGTGTGGCTGGCAATATACCTGCTCCACACAAG
GGGGTTACGATTGGCTTCTCCAAGAATAGAGATTGCTTACGACAGATTGTTTATTGGTCAAGATGGTGTATTACA
AGAATTGTTGCAAGCAAGTTAGAAGCAGAAACTGCTACGAATACCAACAAACATCATGGGAAGAATATGATAGCCAAG
CAGAGAAACGAGTCTATTATTGAGATCAGCGTAGTTACATACACTTAAAGCTGGTGGAGAGCTAGCAGTGGTTGGGTTAAGGA
TATTATTACAGAAGGATGGCTTGTGTTGATTCGCGCATCACAGATTGACGGTTGGAGAGCTAGCAGTGGTTGGGTTAAGGA
TTACCTCTTACGTATGAGAGAGAAGCTAAAGCAGCTCATGGTACTATCTAAATCCAGCAACTGGCATTATGCAACACG
GTGGAATATCTAGGTAATAGATGGTACTACCTCATTGCTCAGGAGCTATGGCAACTGGCTGGTATAAGGAAGGCTCACT
TGGTACTATCTAGATGCTGAAATGGTATGAGAACTGGCTGGCAAAACCTTGGGAAACAAATGGTACTATCTCCGTCATC
AGGAGCTATGCAACTGGTGGTATCAGGAAAGTCAGTGGTACTATCTAAATGCAAGTAATGGAGATATGAAAACAGGCT
GGTCCAAGTCATGGTAACTGGTACTATGCCTATGATTAGCTGCTTAGCTGTTAATACCACAGTAGGTGGTACTACTTA
AACTATAATGGTGAATGGTTAAGTAA

(SEQ. ID. NO. 188)

MKLLKKMMQIALATFFFGLLATNTVFADDSEGWFVQENGRYYKKGDLKETYWRVIDGKYYYFDPLSGEMVVGWQYIPAPHK
GVTIGSPRIETALRPDWFYFGQDGVLQEFVGKQVLEAKTATNTNKHHGEYDSQAERKRVYYFEDQRSYHTLKTGWIYEEGHW
YIQLKDGGFDSRINRLTVGELARGWVKDYLTYDEEKLKAAPWYLNPATGIMQWTGQYLGNRWYLYHSSGAMATGWWYKEGST
WYLYDAENGDMRTGWQNLGNKWYLRSSGAMATGWWYQESSTWYLYNASNGDMKTGWFQVNGNWYAYDSGALAVNTTVGGYLY
NYNGEWRVZ

ID306 (SEQ. ID. NO. 201)

TTGGCTGGTAGATATGGTCTGCTGTTAGTACAGAAGTGAATGGCTCAAACCTTCAACAGTTAAAACATAAGCTACGGT
TGTAGAAAACCACTGAAAGATTAGAGCTACGCTCTGATCAGTCAGTCTGGTTGGGGAATCTAATGGTAAATGGTATTCT
ATGAGTCTGGTAGTGAAGACAGGGTGGTAAAACAGATGGTAAATGGTACTATTGATGACTTAGGTGTATGCAAGACT
GGATTTGTAACATTCTGGTAGCTGGTATTACTTGAGCAATTAGGTCTATGTTACAGGCTGGGAACAGATGGTAGCAG
ATGGTCTACTTGACGGCTCAGGAGCTATGAAGACAGGGTGGTACAAGGAAAATGGCACTGGTATTACCTTGACGAAGCAG
GTATCATGAAGACAGGGTGGTAAAAGTCGGCACACACTGGTACTATGCCACGGTCTAGGAGCTTGGCTGTGAGCACAACA
ACACCAAGATGGTACCGTGTAAATGGTAAATGGTGAATGGTAAACAGGTTAAACTAG

(SEQ. ID. NO. 189)

LAGRYGSAVQCTEVTAASNLSLTVKTKATVVEKPLKDFRASTS DQSGWVESNGKWFYFESDVKTGWVKTDGKWWYLNLDLGVMQT
GFVKFSGSWYLYNSGAMFTGWTGWDGSRWYFDGSGAMKTGWFYKENGTwYLDAGIMKTGWFVGPWYAYGSGALAVSTT
TPDGYRVNGNGEWNZ

ID307 (SEQ. ID. NO. 202)

ATGAAAATTGAAAAAAACTATGCAAGTGGACTGACAGTATTTTCTTGTTGCTAGGGACCAGTACAGTATTCAGA
TGATTCTGAAGGATGGCAGTTGTCAGAAGAAAACGGAAGAACCTACTACAAAAGGGGACCTCAAAGAAAACCTACTGGCGAG
TGATTGATGGTAAGTACTATTATTGATTCTCATCTGGAGAGATGGTGTGGCTGGCAATATATCCCCTTCCATCTAAA
GGTAGTACAATTGGCTTACCCAAATGGTATCAGATTAGAGGTTTCCAAAGTCAGAGTGGTACTACTTCGATAAAAATGG
AGTGCTACAAGAGTTGTTGGTGGAAAACATTAGAGATTTAAACTAAAGACAGTGGTGAAGAAAGTACGGGAAAACGTG
AAGATTCAAGAATTAAGAAGAGAAGCGTTATTATACGAATTACTTTAATCAAATCATTCTTAGAGACAGGTTGGCTT
TATGATCAGTCTAATGGTATTATCTAGTCAAGACGGAAAATTAATGGAGAAAACCTCTGGTGTGAAGACAGTGTGCGGGGTG
GATAAACGATGATTGACTGGTACTACCTAGATCCAACACTGGTATTATGCAAAACAGGTTGCAATATCTAGGTAATAAGT

GGTACTACCTCCGTTCTCAGGAGCAATGGCCACTGGCTGGTATCAGGAAGGTACCACTGGTATTATTTAGACCACCCAAAT
GGCGATATGAAAACAGGGTGGAAAACCTTGGAACAAATGGTACTATCTCGTTCATCAGGAGCTATGGCAACTGGTGGTA
TCAAGATGGTCAACTGGTACTACCTAAATGCAGGTAAATGGAGACATGAAGACAGGGTGGTCCAGGTCAATGGCAACTGGT
ACTATGCTTATAGCTCAGGTGTTGGCAGTGAATACGACCGTAGATGGCTATTCTGTCAACTATAATGGCAATGGGTTCGG
TAA

(SEQ. ID. NO. 190)

MKILKKTMQVGLTVFFFGLLGTSTVFADDSEGWFVQENGRYYKKGDLKETYWRVIDGKYYYFDSLGEVVGWQYIPFPSK
GSTIGPYPNIGIRLEGFPKSEWYYFDKNGVLQEFVGWKTLEIKTKDSVRKYGEKREDSEDKEEKRYYTNYYFNQNHSLETGWL
YDQSNWYLAKEINGENYLGGERRAGWINDDSTWYLDPTTGIMQTGWQYLGNKWYLYRSSGAMATGWYQEGTTWYLDHPN
GDMKTGWQNGLNKWYLYRSSGAMATGWYQDGSTWYLYNAGNGDMKTGFQVNGNWYYAYSSGALAVNTTDGYSVNYGEWVR
Z

ID308 (SEQ. ID. NO. 203)

ATGAAAAAGAATTAACACTGTTAGCACTTGTAGGCCCTTTAGGTTGTATGGTATGGGAATGTCAGGCTCAAGAAAG
TTCAGGAAATAAAATCCACTTTATCAATGTCAGAAGGTGGCAGTGATGCGATTATCTTGAAGCAATGGACATTTGCCA
TGGTGGATACAGGAGAAGATTATGATTCCCAGATGGAAGTGATTCTCGCTATCCATGGAGAGAAGGAATTGAAACGTCTTAT
AACGATGTTCTAACAGACCGTGTCTTCGCTTGAAGGAATTGGGTGCTAAAACCTTGATTTTATTTGGTGAACCCATAC
CCACAGTGATCATATGGAAATGGTGTGAAATTACTGTCACCTATCCAGTGACGAGTCTATCTTAAAGAAATTAAGTG
GTCGTATTACTAATTCTGAACGTCTATGGGATAATCTGTATGGCTATGATAAGGTTTACAGACTGTCAGGAAAGGGTGT
TCAGTTATTCAAATATCACACAAGGGATGCTCATTTGAGTTGGGACATGGATATTCAAGCTCTATAATTGAAATGA
AACTGATTCACTGGGTGAATTAAAGAAAATTGGGATGACAATTCCAATTCTTGATTAGCGTGGTGAAGTCATGGCAAGA
AAATTACCTGGGGCAGTTAGATAATGTCATGGAGCAGAAGACAAGTATGGTCTCTCATGGAAAAGTTGATTGATG
AAGTTAACATCACCATGATAACCAAAATCAACAGGATTGCTTAAAGGAAATTGGAGTTGAGTGTGATTGTTAACTG
TTCGGATAGTCTACCTGGAAAATGGTGTGATGAGTGTGATTGTTAACTGGCTCAAGAACGAGGAATTGAGAGAATCAACG
CAGCCAGCAAAGACTATGCAACAGTTTGATATTGAAAAGACGGTTTGTCAATATTCAACATCTACAAGCCGATT
CCAAGTTTCAAGCTGGTGCATAAGAGTGCATATGGGACTGGTGTATCAAGCGCTGATTCTACAGGAGAGTATGCTGT
CGGTTGGATGAAATCGAAGGTGAATGGTATTACTTAAACCAACGGTATCTTGTACAGAATCAATGGAAAAAATGAAACA
ATCATTGGTTATTGACAGACTCTGGTCTCTGCTAAAGGAAATTGGTGAACGCTGGTAACTGGTATTATTGATGTTCTATG
GAAACAGATGGAATTGGTGGATTCAAGATAAAAGAGCAGTGGTATTATTGGATGTTGATGGTCTATGAAGACAGGATG
GCTTCAATATATGGGCAATGGTATTACTTGCTCCATCAGGGAAATGAAAATGGCTGGTAAAGATAAAAGAAACCTGGT
ACTATATGGATTCTACTGGTGTATGAAGACAGGTGAGATAGAAGTGTGTTCAACATTATCTGAAGATTCAAGGAGCT
ATGAAGCAAGCTGGCATAAAAGGCAATGATGGTATTCTCAAGACAGACGGTTCAGGACTGTGGTTGGATCAAGGA
CAAGGATAATGGTACTCTTGTGAAAGAAAATGGTCAATTACTTGTGAAACGGGACACAGGAAAGGTTACTGTGGATTCA
GTGGTGCTGGTTAGTGGATGTTGATCAGGAAATCTGCTACAATTAAACACTAACAGTCAATTGAGTGTACA
GAAGTAGTGAAGGATCTGAAAATAGAACAGACTGAAAGTGTACAATTTCACTAGTCAAGATTGAC
ATCCTCAACTCACAAAGCTCTGAAACAGAGTGTAAACAAATCGGAATCAGAACAGTAG

(SEQ. ID. NO. 191)

MKKKLTSLALVGAFLGLSWYGNVQAQESSGNKIHFINVQEGGSDAIILESNGHFAMVDTGEDYDFPDGSRSYPWREGIETSY
KHVLTRVFRRLKELGVQKLDFFILVTHSDHIGNVDELLSTYPVDRVYLKKYSDSRITNSERLWDNLGYDKVLOAAEKGV
SVIQNITQGDAHFQFGMDIQLYNENETDSSGELKKIWDDNSNSLISVVVKVNGKIIYLGGLDNVHGAEDKYGPLIGKV
KFNHHHDNTKSNTKDFIKNLPSLIVQTSDSLPLWKNGVDSEYVNLKERIRINAASKDYATVFDIRKDFVNISTSYKPI
PSFQAGWHKSAYGNWWYQADPSTGEYAVGWEIEGEWWYFNQTCILLQNQWKWNHWFYLTDSGASAKNWKIAGIWWYFNK
ENQMEIGWIQDKEWYQYLLDGSMSKMTGWLQYMGQYYFAPSGEMKGMGVWLDKETWYMDSTGVMTGEIEVAGQHYLEDGA
MKQGWHKKANDWYFYKTDGSRAVGWIKDKDWYFLKENGQLLVNGKTPEGYTVDSGAWLVDVSIEKSATIKTTSHSEI
EVVKDLENKETSQHESVTNFSTSQDLTSSTSQSSETSVNKSSEQZ

ID309 (SEQ. ID. NO. 204)

ATGGAATTAACAGATTGAGTAAAGAACAGATTGCTCAAGTCGGCGTCAACCATAAGGAAGTACACGCACACTCAAC
TGGGAATCCGATTCAACCGTACAGAATGAAGCGGATTATCACTGGCGAAAGACCCAGAATTAGGTTTCTCGCACATTG
TTGGGAACGGTTGCATCATGCAAGTAGGACCTGTTGATAATGGTCTGGACGTTGGGAGCTGGGGCGTTGGAATGCTGAGACCTAT
GCAGCGGTTGAACTGATTGAAAGCATTCAACCAAAGAAGAGGTTGACGCGACTACCGCCTTATATCGAATCTTACGCAA
TCTAGCAGATGAAGCAGGTTGCGCAAACAGCTGATACAGGGAGTTAGCTGAAATTAAACACGACAGTATTGACGAA
ACCAACCAAACACACTCAGAACCGTGGACCTTACATCTGCTAAATGGGCTTAGCGTGAGCAGTTAAC
GATATTGAGAACGGCTTGACGATTGAAACAGGCTGGCAGAAGAATGACACTGGCTACTGGTACCTACATTGAGCAGGCTTAA
TCCAAAAGACAAGTTGAGAAAATCAATGGCACTTGTACTACTTGACAGTTGAGCTATATGCTGAGACCGCTGGAGGA
AGCACACAGACGGCAACTGGTACTGGTGCACAACTCAGGCGAAATGGCTACAGGCTGGAGAAAATCGTGATAAGTGGTAC
TATTCTAACGAAGAAGGTGCCATGAAGACAGGCTGGTCAAGTACAAGACACTTGGTACTACTAGCCTAAAGAACGGCG
CATGGTATCAAATGCCATTCTGAGCTCAGCGGACGGAAACAGGCTGGTACTACCTCAAACCAAGACGGAACACTGGCAGACAAGC
CAGAATTCAACAGTAGAGCCAGATGGCTTGATTACAGTAAATAA

(SEQ. ID. NO. 192)

MEINVSKLRTDLPQVGVPYRQVHABSTGNPHSTVQNEADYHWRKDPELGFPSHIVGNICIMQVGPVDNGAWDVGGWNAETY
AAVELIESHSTKEFMTDYRLYIELLRNLADEAGLPKTLTGSLAGIKTHEYCTNNQPNHSDHVDPPYPLAKWGISREQFKH
DIENGLTIETGWQKNDTGWYVHSDGSYPKDKFEKINGTWYLYFDSSGMLADRWRKHTDGNWYWDNSGEMATGWKKIADK
WYFNEEGAMKTGWVVKYKDTWYLLDAKEGAMVSNAFIQSADGTGWYLYLKPDTLADKPEFTVEPDGLITVKZ

ID310 (SEQ. ID. NO. 205)

ATGGCACAACAGGATTACAATAATTGACTTAATTATCTTGATTGTTATTTACTTGCAGGTGTTGGTGCAGGTATCTATT
CTCTAAAAAGAGATGAAAGAAAAGAGTCTTAAAGGAGATGGTCTGGTCTGGTATGTTACTTCGGTATCCATT
CCACAATGCTCAGTCCGATTCTCTGGACTCGCTGGTAGCTTATGCAGGTAGCTGGATTGGTTGCTCAATT
GGGATGGTAGTGTGCTTACCTGACAAATTGTTTATCTTACCTATCTTGACAGGATAGACATCGATAACGGCATATGATTA
CTTGGATAACGTTTAACTCTAACAGACTCTGTTATCTGAGTACAGGCTGGTCAATTGAGACACTTGGTACTACT
TCATTATGACCTCCCATGCTGGTATCTGAGTACAGGAGTACATCAATTGGTATTGATTTGATGGGTGAGT
GCAATTGTTATTCTTACTGGTGGCTAAACCGTATTATGGACAGACTTATTCAAGGTGCTGATTAGTGGTGT

CGTTTAGCTTATTTGACTGATTGCTAATATTAAGGGCTTGGCAGTAGCAGAACATTAGCAAACGGAAATTCC
TTGCTGCAAATGAAAACCTTCGATCTAACTGCTTCAAACTCCATCTTTAATTGTGATGGGTCAGGCTTACAATC
TTGCTTCCCTATGCTCATCTCAAGATTGGTCAACGTTTACTACAACACAAAATATTAAGAAACTTAAGATGTTGTT
CAAAACGGTGTGTTGCACTGCAACTGCAACAGCTTTACTGATGGTACAGGCTTGACGTATTCTATCAAGTACAAA
ATGCAGATAGTGCAGCTGCAATCTCCCTCAAGACCAAATCTTATGACTTTACTGCATACCAGTTACAGTAGGTATCACA
GGTTGATCTGGCAGCGATTATGCAGCATCTCAATCAACTATTCACAGGTTGACTCTGTCAGACTCATGGACATT
GGATATTCAAGATGTCATTCAAAAATATGTCAGACAATCGTCAGAAAATTGCAACAATTGCTATCTAGCAGTAGGTT
TATTCTCAATTGGTGTGTTCCATTGTCATGGCTCACTCAGATATTAACCTGCATACGAATGGTCAATAGTTCATGGGACTT
GTACTGGTCACTGGTGTGTTCTTGGATTGTTCTAAAAAGCAAATAAACAGGTGTTAGCTACTGGGCAATTGATTTCAA
TGATCAACCCTGTCATGGTATTATTAACCTCTCCACAGCTGTTAGCTACTGGGCAATTGATTTCAA
TCTCTGATTCAGTAGTTCAAGGTTATGTCATCTGTCAGGTTACTGGGAAATTGATCTGCACCTAAATACAAACGATTCA
GATATTACAGAAATTAAACGGGATTCAGTTGGGAAAGTTCGTCACTAA

(SEQ. ID. NO. 193)

MGGTGTIDLIILIVYLLAVLVAGIYFSKEMKGKEFFKGDSVPWVTSVI FAMLSPISFLGLAGSSYAGSWILWFAOL
GMVVAIPLTIRFILPIFARIDIDTAYDYLDRFNSKALRI I SALLFI I YQLGRMS I IMYLPSSAQLSVLTGIDINILIILMGVV
AIVSYTGGLKSVLWTDFIQGVILISGVVLALFVLIANIKGGFGAVAETLANGKFLAANEKLFDPNLLSNSIFLIVMGSFTI
LSSYASSQDLVQRFTTQNICKLNKMLFTNGVLSLATATVFLYIGTGLYVFYQVNADSAASNPQDQIFMFYIAYQLPVGIT
GLILAAIYAAQSSTISTGLNSVATSWTLIDQDVISKNMRSRRTKIAQFVSLAVGLFSIGVSIVMAHSDIKSAYEFNSFMGL
VLGLLGGVFIIFGVSKANKQGAYAALIVSTIVMFVFIKYFLPPTAVSYWAYSLISISVSVVSGYIVSVLGNKVSAPKTTIH
DITEIKADSSWEVRHZ

ID311 (SEQ. ID. NO. 206)

ATGAAAATTAATAAAAAATCTAGCAGGTTCACTGGCAGTCCTGCCCTAAGTGTGTTCTATGAGCTGGTCGTACCA
AGCTGGTCAAGGATAAGAAGAGCTAACTGAGTTGCTTATAGATGGTCACTGGCTGGTCAAAGGAGAAAACCTGGACAC
CAGATGAGTCAGTAAGAGGGAGGGATCAACGCCAACAAATCGTCATCAAGATTACGGATCAAGGTTATGTGACCTCTCAT
GGAGACCATTATCATTACTATAATGCAAGGTCCCTATGATGCCATCATCAGTGAAGAGCTCCTCATGAAAGATCCGAAATTA
TCAGTTGAAGGATTAGACATTGCAATGAAATCAAGGGTGGTTATGTCATCAAGGTAGACGGAAAATACTATGTTACCTTA
AGGATGAGCTCATGGCATAATTGGACAAAAGAAGAGATTAAACGTCAGAAGCAGGAACGCAGTCATAATCACGGGTC
GGAGCTAACGATGTCAGTGCAGGGCAGAGGCCAACGGACGATGGTATATCTCAATGCACTGATCTGA
TATCATTGAGGACACGGGTAGCTGCTTATGTCCTCATGGCAGGACATTACCTACATCCCTAAGAATGAGTTATCAGCTA
GCGAGTTAGCTGCTGAGAACGCTATTGGAAATGGAAGCAGGGATCTGCTCTTCTCAAGGTTAGTTATAATGCAAATCCA
GCTCAACCAAGATTGTCAGAGAACACAATCTGACTGCACTCAACTTATCATAAAATCAAGGGAAAATCTCAAGCCT
TTTACGTAATTGATGCTAAACCCCTTATCAGAACGCCATGTGAATCTGATGCCCTTATTTGACCCAGCGCAAATCAA
GTCGAACCGCAGAGGTAGCTGTCCTCATGGTAAACCTACACTTACCTTATGAAACAAATGTCGAAATTGGAAAAAA
CGAATTGCTCGTATTATCCCTCGTATCGTCAAACATTGGTACAGATTCAAGAACGAGAACACAACGTCACAAATC
GACTCCGAAACCTAGTCAAAGTCCGAACTGCACCAATCCTCAACAGCTCAGAACATTGATGAGAAATTGGTCA
AAGAAGCTGTTGAAAAGTAGCGATGGTTATGCTTGGAGGAGATGGAGTTCTGTTATATCCCAGCCAAGGATCTTCA
GCAGAACACAGCAGCAGGATTGATGCAAACACTGCCAAGCAGGAAAGTTATCTCATAGCTAGGAGCTAAGAAAACCTGACCT
CCCATCTAGTATGCAAGAATTTCACAATAAGGTTATGACTACTAGCAAGAATTACCAAGAATTACTGATAATAAAAGGTC
GACAAGTTGATTGAGGCTTGGATAACCTGTGGAAAGCACTGAGGATGTCCTCAAGTGTGATAAAAGTCAAGTTAGTGGATGAT
ATTCTGCCTCTTAGCTCGATTGTCATCCAGAACGTTAGGAAAACCAAATGCGCAAATTACCTACACTGATGATGAGAT
TCAAGTAGCCAAGTTGGCAGGCAAGTACACAACAGACGGTTATCTTGTGATCTCGTATAACCAGTGTGAGGGGG
ATGCCATGTAACCTCCACATGACCCATAGGCCACTGGATTAAAAAGATAGTTGCTGAGCTGAGAGAGCGGCAGCCAG
GCTTATGCTAAAGAGAAAGGTTGACCCCTCTGCAAGACCATCAGGATTGAGGAAACTCAGGAGGAGGAGCAGAAC
TATCACAACCGCGTGAAGCAGCTAAGAAGGTGCACTGTGATCGTATGCTTACATCTTACATATACTGAGTAAAGTCAAG
ACGGTAGTTAATCATACCTCATTGACCAATTACATAACATCAAATTGAGTGGTTGACGAAGGCTTATGAGGCACT
AAGGGGTATACTTGTGAGGATCTTGGCAGTGTCAAGTACTATGTCGAACATCCAACAGTCAGCAGATAATGG
TTTGGTAACGCTAGCGACCATGTTCAAAGAAACAAAATGGTCAAGCTGATACCAATCAAACGGAAAACCAAGCGAGGAGA
AACCTCAGACAAAAACCTGAGGAGAAAGAACCCCTCGAGAGAGAGAACCGCAAAGCGAGAACAGCTCCAAAACCA
GAGGAACAGAAGAATCACCAGAGGAATCAGAAGAACCTCAGGTCAGAGCTGAAAGGTTGAGGAAAATCTGAGAGGGCTGA
AGATTTACTGGAAAATCCAGGATCCAATTATCAAGTCAAAGGACTCTCACAGGATTTAAATTACTAT
TTGGCACCCAGGACAACAAACTATTATGGCAGAAGCTGAAAATCTTGGCTTATTAAAGGAGAGTAAGTAA

(SEQ. ID. NO. 194)

MKINKYLAGSVAVLALSVCSYELGRHQAGQDKESNRVAYIDGDQAGQKAENLTPDEVSREGINAEQIVIKITDQGYVTSH
GDHYHYYNGKVPDAI I SEELLMDPNYQLKDSIVNEIKGGYVKVDGKYYVLDAAHADNIRTKEEIKROKOERSHNHGS
GANDHAVAAARAQGRYTTDDGYI FN ASD I IEDT GDAY I VPHGDHYH I PKNELSAE LAAA EAY WNGKQGSRPSSSSYNANP
AQPRLENHNLTVPPTYHQNQGENISLRLREYAKPLSERHVESDGLI FDPAQITSRTARGVAVPHGNHYHFIPYEQMSELEK
RIARI I PLRYRSNHWVPDSPRPEPSPSPQAPNPQAPSPNPIDEKLVKEAVRKVGDGYVFEENGVSRY I PAK DLS
AETAAGIDSLSKLAQKESLHSLKGAKKTDLPSSDREFYNKVAYDLLARIHQDLDNKGQVDFEALDNLLERLKDVPSDKVVLVDD
I LAFLAPIRHPERLGKPNQAITYTDDEI QVAKLAGKYTEDGYI FDPRDITSDEGDAYVTPHMTHSHWIKKDSLSEAERAAQ
AYAKEKGTLPPSTDHQDSGNTEAKGAEAI YNRVKAACKVPLDRMPYLNQYTVEVKNGSLI I PHYDHYNIKFEWFDEGLYEAP
KGYLEDLLATVKYYVEHPNERPHSDNGFCNASDHVQRNKGQADTNQTEKPSEEKPQTEKPEEETPREEKPKSEPKPESPKPT
EEPEESPEESEEPQVETEKVEEKLREAEDLLGKIQDPIIKSNAKETLTGLKNLLFGTQDNNTIMAEEAKLALLKESKZ

ID312 (SEQ. ID. NO. 207)

ATGGAGGGATTGGTTAGAGTCATTATGCCGTATTTGGCATTACAAGCTATCTAAACCTACTACGCCATTCTTCAACA
GCAAGTAAACAAATGGGCTGACAAGGCAAATAAGGCGAAAAGGGGCAATTGCTAACTACTCTTGTCCATAACATGAATA
AGCGTATTGAAATATGCCGTAGCTATCCAGGTAACTACAACCCAGCTATGATGTCATCGTCCACGCCAACAGCAA
AAAGAAAAGGCTGCTGCAAATACTTAGACAACAAAGAATTAAACAGTTCTGATTATTTAGATGCTCTGGATCAATCAA
TTATGAGAACTTATGATGTTCTGTATAAGACTTTATGCCACTGGTTGCCATTAGTGAGGCTCTGGCTTCA
GGTCTGATATTGACCTAGAAAGCGGTGTTATCAGCATCAATAAGACACTAAACCGCTATCAGGAAATAACCTCACCTAAATCA
AGCGTGGTTATCGTGTATACCAATAGACAACAGCCACATTACTTTACTGAAACAATACAAAACCGTCAACAAATTGAGTC
TTGAAATTAGGCCGATCTGAAACAGTTGATTCTCTGTATTACGGAGAAATATGCTTATGCTGTAACCTACGCCAACAGCC
TAAATAAGCATTGATGCTGGAGTAACAGTATCTCATGGTTCCGCCATACACATACTACTATGATGCTTAT

GCTCAGGTTAGCCCGAAAGATGTTCAGTATAGATTAGGCCACTCTAATTAAATGATCACTGAAAATACCTACTGGCATACTAA
CCAAGAGAATGAAAAAAAGCCGTCTCAAATTATGAAACAGCTATCAACAATTATAA

(SEQ. ID. NO. 195)

MEGLVRVHLLPVFGDYKLSKLTTPILQQQVNWKADKANKGEKGAFANYSLLHNMMNKRILKYGVAIQVIQYNPANDVIVPRKQQ
KEKAALKYLDNKELKQFLDYLDALDQSNYENLFDVVLYKTLLATGCRISEALALEWSDIDLESGVISINKTLNRYQEINSPKS
SAGYRDIPIDKATLLLLKQYKNRQQIQSWKLGRSETVVFSVFTEKYAYACNLRKRLNKHFDAAGVTNVSFHGFRHTTMMLY
AQVSPKDVQYRLGHSNLIMITENTYWHTNQENAKKAVSNYETAINNLZ

At Page 1, 2nd full paragraph:

Background of the Invention

Streptococcus pneumoniae, commonly referred to as the pneumococcus, is an important pathogenic organism. The continuing significance of *Streptococcus pneumoniae* infections in relation to human disease in developing and developed countries has been authoritatively reviewed (Fiber, G.R., *Science*, 265:1385-1387 (1994)). That indicates that on a global scale this organism is believed to be the most common bacterial cause of acute respiratory infections, and is estimated to result in 1 million childhood deaths each year, mostly in developing countries (Stansfield, S.K., *Pediatr. Infect. Dis.*, 6:622 (1987)). In the USA it has been suggested (Breiman *et al*, *Arch. Intern. Med.*, 150:1401 (1990)) that the pneumococcus is still the most common cause of bacterial pneumoniae, and that disease rates are particularly high in young children, in the elderly, and in patients with predisposing conditions such as asplenia, heart, lung, and kidney disease, diabetes, alcoholism, or with immunosuppressive disorders, especially AIDS. These groups are at higher risk of pneumococcal septicaemia and hence meningitis and therefore have a greater risk of dying from pneumococcal infection. The pneumococcus is also the leading cause of otitis media and sinusitis, which remain prevalent infections in children in developed countries, and which incur substantial costs.

At Page 3, 2nd full paragraph:

Brief Summary of the Invention

Thus, in a first aspect the present invention provides a *Streptococcus pneumoniae* protein or polypeptide having a sequence selected from those shown in Table 2 herein.

At Page 3, 2nd full paragraph:

Detailed Description of the Invention

As discussed herein, the proteins and polypeptides of the invention are useful as antigenic material. Such material can be "antigenic" and/or "immunogenic". Generally, "antigenic" is taken to mean that the protein or polypeptide is capable of being used to raise antibodies or indeed is capable of inducing an antibody response in a subject. "Immunogenic" is taken to mean that the protein or polypeptide is capable of eliciting a protective immune response in a subject. Thus, in the latter case, the protein or polypeptide may be capable of not only generating an antibody response and in addition non-antibody based immune responses.

At page 101, line 1:

CLAIMS *What is claimed is:*